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RESULT 20
GRAR RANRU
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AC 940754
DT 01-FEB
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DE Granul
OS Rana r
OC Amphib
OX NCBI_T
RN [1]
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RC TISSUE
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RI "A new
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KW Amphib
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=78189201; PubMed=657408;
Nakajima T., Yasuhara T., Hirai Y., Kitada C.,
Koyama K., Yajima H.;
"Synthesis of the dodecapeptide amide correspor
amino acid sequence of granuliberin-R, a new fx
Rana rugosa.";
Chem. Pharm. Bull. 26:1222-1230(1978).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nakajima T., Yasuhara T.;
"A new mast cell degranulating peptide,
(Rana rugosa) skin.";
Chem. Pharm. Bull. 25:2464-2465(1977).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rana rugosa (Wrinkled frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-FEB-1995 (Rel. 31,
01-FEB-1995 (Rel. 31,
10-OCT-2003 (Rel. 42,
Granuliberin-R.
                                                                                                                                                                                                                                                                                                                                                                              NP1_LYMST STANDARD; PRT; 13 AA P80178; 01-JUL-1993 (Rel. 26, Created) 01-JUL-1993 (Rel. 26, Last sequence update) 01-JUL-1993 (Rel. 26, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amphibian defense peptide; Mast cell degranulation; Amidation.

MOD_RES 12 12 AMIDATION.

SEQUENCE 12 AA; 1424 MW; 2B974EB9CA1B5047 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=8410;
MEDLINE=93238777; PubMed=8477756;

Johnsen A.H., Rehfeld J.F.;

"LymnaDFamides, a new family of neuropeptides from the pond Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity invertebrates?";

Eur. J. Biochem. 213:875-879(1993).

-i- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
                                                                                                                                                                                                                                                                            Ol-UULTION
Lymna-DF-amide 1.
Lymnaea stagnalis (Great pond snail).
Lymnaea stagnalis (Great pond snail).
Lymnaea stagnalis (Great pond snail).
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MEDLINE=78062810; PubMed=589733;
                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                   sukaryota; Metazoa; Mollusca; Gas
symnaeoidea; Lymnaeidae; Lymnaea.
ICBI_TaxID=6523;
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SUBCELLULAR LOCATION: Se
TISSUE SPECIFICITY: Skin
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Pred. No. 5.5e
1; Mismatches
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5.5e+03;
hes 0;
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Ranidae; Rana
                                                                                                                                                                                                                                                                                                                     Basommatophora;
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RESULT 22
NP4_LYMST
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MOD_RES 1:
SEQUENCE 13 1
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P80181;
01-JUL-1993
01-JUL-1993
01-JUL-1993
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Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
Lymnaeoidea; Lymnaeidae; Lymnaea.
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                                                                                                                                                                                                             Neuropeptide; Amidation.
MOD_RES 13 13
SEQUENCE 13 AA; 1503 N
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MEDLINE=93238777; PubMed=8477756;
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                                                                                                                                                                                                                                                                                               -!- SIMILARITY: RELATED
PIR; S32474; S32474.
                                                                                                                                                                                                                                                                                                                                                                                             Johnsen A.H., Renfeld J.F.;
"LymnaDFamides, a new family of neuropeptides from the pond Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity
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                                                 6 NSKF 9
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NSAF 10
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(Rel. 26, Last sequence up)
(Rel. 26, Last annotation
                                                                                                           Conservative
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RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
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75.08;
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9CA07BA3F5D5B455
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9CA07BBB56D5B455 CRC64;
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5.9e+03;
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RESULT 23
TEML_RANTE
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AC P57104;
      CCCL 11 PAXCO CCC CCC TANK CCC CCC CCC TANK CCC TANK CCC CCC TANK CCC TANK CCC TANK CCC TANK CCC CCC TANK CCC TANK CCC CCC TANK CCC
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16-OCT-2001
10-OCT-2003
Temporin L.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea;
NCBI TaxID=8407;
                                                                                                                                                                                           Simmaco M., Mignogna G., Barra D.;
                                                                                                                                                                                                                                                                    TISSUE=Skin secretion; MEDLINE=97175050; PubMed=9022710;
                                                                                                               temporaria.
                                                                                                                                                           "Temporins,
                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                         J. Biochem. 242:788-792(1996) FUNCTION: Has antibacterial ac
Gram-positive
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(Rel.
(Rel.
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                                            activity against Gram-negative
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Ranidae; Rana
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frog

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RESULT 25
RS19_PPWBP
ID RS19_P
AC Q52093
AC Q52093
DT 30-MAY
DT 30-MAY
DT 30-MAY
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Matches 3
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MEDLINE=94350802; PubMed=8071198;
Gundersen D.E., Lee I.M., Rehner S.A.,
"Phylogeny of mycoplasmalike organisms
their classification.";
J. Bacteriol. 176:5244-5254(1994);
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Ukena K., C
                                                                                               Pigeon pea witches'-broom phytoplasma Bacteria; Firmicutes; Mollicutes; Ach
                                                                            Acholeplasmataceae; Phytoplasma.
NCBI_TaxID=37700;
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-!- TISSUE SPECIFICITY: Skin.
-!- SIMILARITY: Belongs to the brevinir Amphibian defense peptide; Antibiotic; MOD RES 13 13 AMIDATION.
SEQUENCE 13 AA; 1641 MW; 9EBDCB1FAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eisenia foetida (Common brandling worm) (Common dung-worm)
Eukaryota; Metazoa; Annelida; Clitellata; Oligochaeta; Hap
                                                                                                                                                                                                                                                                                                                                           Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                   Peptides 16:995-999(1995).
-!- FUNCTION: Has a stimulative effect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation updat
Myoactive tetradecapeptide (ETP)
                                                                                                                                                                                                                                                                                                                                                                                                           foetida."
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Ukena K., Oumi T., Matsushima O., Ikeda T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lumbricina;
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PSS OR RPS19
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Rel. 41, Last annotation updat
protein S19 (Fragment)
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44
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Pred. No. 6.3e
2; Mismatches
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CC9ABEF941CD91AD
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Pred. No.
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                     Davis R.E., Kingsbury D. (phytoplasmas): a basis
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RESULT 26
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Best Local S
Matches 3
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P20750;
01-FEB-1991
01-FEB-1991
16-OCT-2001
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-i- я
           Conlon J.M., Deacon C.F., Hazon N., Henderson I.W., "Somatostatin-related and glucagon-related peptides structural features from the European eel (Anguilla Gen. Comp. Endocrinol. 72:181-189(1988).

-i- FUNCTION: Somatostatin inhibits the release of s
                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Crama Actinopterygii; Neopterygii; Teleos Acanthomorpha; Acanthopterygii; Per Cottoidei; Cottidae; Myoxocephalus. NCBI_TaxID=8097, 8019, 7936;
                                                                                                                                                                         SPECIES=O.kisutcn; 115000-17919;
MEDLINE=87055212; PubMed=2877919;
Plisetskaya E.M., Pollock H.G., Rov
Plisetskaya Andrews P.C., Gorbman,
                                                                                                                                                                                                                                                         Conlon J.M., Davis M.S., Falkmer S., Thim L.; "Structural characterization of peptides derived from prosomatostatins I and II isolated from the pancreatic islets o species of teleostean fish: the daddy sculpin and the flounder. J. Biochem. 168:647-652(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                Myoxocephalus scorpius (Shorthorn sculpin) (Oncorhynchus kisutch (Coho salmon), and Anguilla anguilla (European freshwater eel).
                                                                                SPECIES=A.anguilla; TISSUE=Pancreas; MEDLINE=89065329; PubMed=2904391;
                                                                                                                                                               "Characterization of coho
                                                                                                                                                                                                                                                                                                                             SPECIES=M.scorpius;
MEDLINE=88029486; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Somatostatin I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ribosomal protein; rRNA-binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    between the Swiss Institute of Bioinformatiche European Bioinformatics Institute. They use by non-profit institutions as long a modified and this statement is not removed. entities requires a license agreement (See ) or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                             SPECIES=M
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InterPro; IPR002222; Ribosomal_S19.
PROSITE; PS00323; RIBOSOMAL_S19; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -!- FUNCTION: Protein S19 forms a complex with S13 that binds strongly
to the 16S ribosomal RNA (By similarity).
-!- SIMILARITY: Belongs to the S19P family of ribosomal proteins.
                                                                                                            SEQUENCE.
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between the Swiss Institute of Bioinformatics and the EMBL
. Comp. Endocrinol. FUNCTION: Somatosta SUBCELLULAR LOCATION
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(Rel.
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. 17, Last seq
. 40, Last anno
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PubMed=2889597;
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75.0%;
                                                                                                                                     63:252-263(1986).
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                                                                                                                                                                salmon (Oncorhynchus
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Pred. No.
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                                                                                                                                                                                         Rouse J.B., Hamilton J.W.,
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RESULT 28
LCK5_LEUMA
ID LCK5_LEUMA
AC P19987;
DT 01-FEB-1991
DT 01-FEB-1991
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Best Local S
Matches 3
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SEQUENCE.
SPECIES=A.mississippiensis; TISSUE=Stomach;
SPECIES=A.mississippiensis; TISSUE=Stomach;
SPECIES=A.mississippiensis; TISSUE=Stomach;
MEDLINE=93324451; PubMed=8101369;
Wang Y., Conlon J.M.;
Wang Y., Conlon J.M.;
"Neuroendocrine peptides (NPY, GRP, VIP, some stomach of the alligator.";
Peptides 14:573-579(1993).
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P31885;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES=T.scripta;
MEDLINE=90341082; PubMed=1974347;
Conlon J.M., Hicks J.W.;
"Isolation and structural characterization of insulin, somatostatin from the turtle, Pseudemys scripta.";
Peptides 11:461-466(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alligator mississippiensis (American alligator), and Trachemys scripta (Red-eared slider turtle) (Pseudemys scripta). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Crocodylidae; Alligatorinae; Alligator.
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Pfam; PF03002; Somatostatin; 1.
Hormone; Multigene family.
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01-JUL-1993 (Rel. 26, Last seq
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!- SIMILARITY: Belongs to the soma
PIR; C60414; C60414.
InterPro; IPR004250; Somatostatin.
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3; Conserv
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sequence
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6.3e+03;
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RESULT 29
SAMP_MUSCA
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P19095;
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01-NOV-1990
28-FEB-2003
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MEDLINE=83160932; PubMed=6403520;

Robey F.A., Tanaka T., Liu T.-Y.;

Robey F.A., Tanaka T., Liu T.-Y.;

"Isolation and characterization of two major
                                                                                          Lectin; A
DOMAIN
NON TER
SEQUENCE
                                                                                                                                                                                                                                                 component.";
J. Biol. Chem. 258:3889-3894(1983).
-!- SUBUNIT: Homopentamer. Pentaxin (or penarrangement of 5 noncovalently bound su-
-!- SUBCELLULAR LOCATION: Secreted.
-!- SUBCELLULAR LOCATION: METERIA OF AMYLOID
-!- DISEASE: SAP IS A PRECURSOR OF AMYLOID
IN BASEMENT MEMBRANE AND ASSOCIATED WIT
-!- SIMILARITY: Belongs to the pentaxin fam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=87052651; PubMed=2877794;
Holman G.M., Cook B.J., Nachman R.J.;
"Isolation, primary structure, and synthesis of
myotropic peptides of Leucophaea maderae.";
Comp. Biochem. Physiol. 88C:27-30(1987).
-i- FUNCTION: This cephalomyotropic peptide sti
activity of cockroach protodeum (hindgut).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blaberoidea; Blaberidae; Leucophaea.
                                                                                                                                                                                                                                    PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mustelus.
NCBI_TaxID=7812;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes; Triakidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-OCT-2003 (Rel.
Leucokinin V (L-V)
                                                                                                                                                                                   PROSITE;
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-!- SUBCELLULAR LOCATION:
PIR; JS0315; JS0315.
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TISSUE=Head;
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                                                                                                                                                                                                                                    B20569;
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p 27.5%;
Similarity 44.4%;
4; Conservative
                                                                                                                                         Amyloid; Glycoprotein; Plasma; 1 >9 PENTAXIN.
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id P-component (SAP) (Fragment).
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PENTAXIN; PARTIAL.
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33.3%;
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736365A5B9C865B8
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                                                                                              D05B5735B3386769
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2; Mismatches
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ID FAR2 PENMO
AC P83317;
DT 28-FEB-2003 (
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Matches 2
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01-DEC-1992
01-DEC-1992
01-NOV-1995
                                                                                                                                            gene sequences.";
J. Bacteriol. 174:2606-2611(1992).
-!- FUNCTION: This protein binds directly to 23S ribo
located at the a site of the peptidyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Seven novel FMRFamide-like neuropeptide sequences from of the giant tiger prawn Penaeus monodon."; Comp. Biochem. Physiol. 131B:325-337(2002).
-!- SUBCELLULAR LOCATION: Secreted.
-!- MASS SPECTROMETRY: MW=1260.0; METHOD=MALDI.
-!- SIMILARITY: Belongs to the FARP (FMRFamide related p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
FMRFamide-like neuropeptide FLP2 (AYSNLNYLRF-amide).
Penaeus monodon (Penoeid shrimp).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
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MOD_RES 10 10
SEQUENCE 10 AA; 1260
                         the
                                                                                                       similarity).
-!- SIMILARITY: Belongs to the L16P family of ribosomal proteins.
                                                                                                                                                                                                                                                 organism and Acholeplasma
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MEDLINE=92210505; PubMed=1556079;
Lim P.O., Sears B.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Firmicutes; Mollicutes; Acholeplasmatales;
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Sithigorngul P., Pupuem J., Krungk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eumalacostraca; Eucarida; Decapoda;
Penaeidae; Penaeus.
                                              between
                                                                                                                                                                                                                                                 "Evolutionary relationships of a plant-pathogenic mycoplasmalike organism and Acholeplasma laidlawii deduced from two ribosomal p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chaivisuthangkura P.,
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Bioinformatics Institute.
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Sithigorngul W., Petsom
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DT 15-JUL-1998 (
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9 CLYEP STANDARD Q46490; 30-MAY-2000 (Rel. 39, C 30-MAY-2000 (Rel. 39, L 28-FEB-2003 (Rel. 41, L 30S ribosomal protein S RPSS OR RPS19.
                                                                                                                                                                                                                                                                                                        "Two-dimensional gel electrophoresis separation and N-terminal sequence analysis of proteins from Clostridium pasteurianum W5 Electrophoresis 19:802-806(1998).
                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                              Elongation NON_TER
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                                                                                                                                                                                                                                                                                                                                                             STRAIN=W5;
MEDLINE=98291870; PubMed=9629918;
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                                                                                                                                                                                                                                InterPro; IPR000795; EF GTPbind: PROSITE; PS00301; EFACTOR GTP; F
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PROSITE; PS00586; RIBOSOMAL_L16_1;
PROSITE; PS00701; RIBOSOMAL_L16_2;
                                                                                                                                                                                                                                                                                                                                                                                                                  Clostridium.
                                                                                                                                                                                                                                                                                                                                                   Flengsrud R., Skjeldal
                                                                                                                                                                                                                                                                                                                                                                                   EQUENCE
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EF-G/EF-2 subfamil
                                                                                                                                                                                                                                                                            SUBCELLULAR LOCATION:
                                                                                                                                                                                                                                                                                        ribosome.
                                                                                                                                                                                                                                                                                               the nascent protein chain from
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3; Conserv
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98 (Rel. 36, Last sequ
03 (Rel. 41, Last anno
n factor G (EF-G) (CP
                                                                                                                                                                                                    11 AA;
                                                                                                                                                                                                                       S00301; EFACTOR GTP; PARTIAL. factor; Protein biosynthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10 AA;
          (Rel. 39, Created)
(Rel. 39, Last sequence update)
(Rel. 41, Last annotation update)
il protein S19 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                    pasteurianum
                                                                                                                                                            Conservative
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50.0%;
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                                                                                                                                                                                                                                                                Cytoplasmic.
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(See http://www.isb-sib.ch/announce/
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RESULT 34.

RS19_TOBBP
ID RS19 T
AC Q56251
DT 30-MAY
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DE 30S ri
GN ROBATO
OC Bacter
OC Achole
OX NCBI_T
RN [1]
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CC -i- FU
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Best Local S
Matches 3
                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no wa modified and this statement is not removed. Usage by and for commercia entities requires a license agreement (See http://www.isb-sib.ch/announce or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; L27019; AAA83940.1; -.

HAMAP; MF 00531; -; 1.

InterPro; IPR002222; Ribosomal S19.

PROSITE; PS00323; RIBOSOMAL S19; PARTIAL.

Ribosomal protein; rRNA-binding.

NON TER 1 1

NON TER 1 1

SEQUENCE 12 AA; 1409 MW; 4CC478EFC833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gundersen D.E., Lee I.M., Rehner S.A., Davis R.E., Kingsbury D.T.; "Phylogeny of mycoplasmalike organisms (phytoplasmas): a basis for their classification.";
J. Bacteriol. 176:5244-5254(1994).
-!- FUNCTION: Protein S19 forms a complex with S13 that binds strongly to the 16S ribosomal RNA (By similarity).
-!- SIMILARITY: Belongs to the S19P family of ribosomal proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Firmicutes; Acholeplasmataceae; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Clover yellow edge phytoplasma.
Bacteria; Firmicutes; Mollicutes; Acholeplasmatales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. MEDLINE=94350802; PubMed=8071198;
                                                                                                                                                                                          J. Bacteriol. 176:5244-5254(1994).
-!- FUNCTION: Protein S19 forms a complex with S13 that binds str
to the 16S ribosomal RNA (By similarity).
-!- SIMILARITY: Belongs to the S19P family of ribosomal proteins.
                                                                                                                                                                                                                                                                  MEDLINE=94350802; PubMed=8071198; Gundersen D.E., Lee I.M., Rehner "Phylogeny of mycoplasmalike orgather classification.";
                                                                                                                                                                                                                                                                                                                                                                                                                           Tomato big bud phytoplasma.
Bacteria; Firmicutes; Mollicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RPSS OR RPS19
                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=35770;
                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAY-2000
28-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                         choleplasmataceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ribosomal
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      L27004;
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3; Conserv
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(Rel. 39, Last sequence update)
(Rel. 41, Last annotation update)
il protein S19 (Fragment).
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      AAA83952.1;
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                                                                                                                                                                                                                                                                                                                                                                                                        Phytoplasma.
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organisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4CC478EFC8333AA3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                              Acholeplasmatales;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8.4e+03
                                                                                                                                                                                                                                                                                        Davis R.E., Kingsbury D. (phytoplasmas): a basis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
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Best Local :
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01-FEB-1991
01-FEB-1991
                                                                                  NON_TER
ACT_SITE
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HAMAP; MF_00531; -; 1.
InterPro; IPR002222; Ribosomal
PROSITE; PS00323; RIBOSOMAL_S1
                                                                                                                                                                                                                                                                                        Bailey J.M., Colman R.F.;
"Isolation of the glutamyl peptide labeled by the nucleotide analogue
2-(4-bromo-2,3-dioxobutylthio)-1,N(6)-ethenoadenosine
2',5'-biphosphate in the active site of NADP+-specific isocitrate
                                                                                                                                                                                                                                                                                                                                                                                                                                           O1-FEB-1991 (Rel. 17, Created)
O1-FEB-1991 (Rel. 17, Last sequence update)
O1-OCT-1996 (Rel. 34, Last annotation update)
Isocitrate dehydrogenase [NADP] cytoplasmic (EC 1.1.1.42)
(Oxalosuccinate decarboxylase) (IDH) (NADP+-specific ICDH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribosomal protein; rRNA-binding.
                                                                                                                           InterPro; IPR001804; Isodh.
PROSITE; PS00470; IDH_IMDH; PARTIAL.
Oxidoreductase; NADP; Tricarboxylic
                                                                                                                                                                              dependent isocitrate dehydrogenase.";
J. Biol. Chem. 262:12614-12619(1987).
-!- CATALYTIC ACTIVITY: Isocitrate + 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                               -!- SUBUNIT: Homodimer.
InterPro; IPR001804; Isodh.
                                                                                                                                                                                                               "Characterization of an active site analogue 3-bromo-2-ketoglutarate on
                                                                                                                                                                                                                                               MEDLINE=87308292; PubMed=3624273;
                                                                                                                                                                                                                                                                                                                                               MEDLINE=87308293; PubMed=2887570;
                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                              Sus scrofa
                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                 (Fragment).
                                                                                                                                                                                                                                                                                Biol.
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                                        Similarity
3; Conserv
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                    GYKNSK 8
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GLSNVK
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                                         Conservative
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                                                   27.5%;
50.0%;
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Т.
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                                                                                                                                                                                 Isocitrate + NADP(+) = 2-oxoglutarate + CO(2
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_S19; PARTIAL
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                                         0
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                                                             Score 14;
                                                    Pred.
                                                                                  1B640F0E9F7C71E0 CRC64;
                                                                                                         POTENTIAL
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                                         Mismatches
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a single chain of
                                                                                                                             acid
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                                                    9e+03;
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                                                                                                                             cycle
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                                                             Length 13;
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f dimeric NADP+-
                                                                                                                                                                                                                                                                                                                                                                                                                                              (IDP)
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RESULT 36
NEUT CAVPO
ID NEUT CAVPO
AC P32560;
DT 01-OCT-1993

STANDARD;

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P32560; 01-OCT-1993

(Rel. 27,

Created)

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RESULT 37

UC18_MAIZE

ID UC18_MA

AC P80624

DT 01-OCT

DT 01-OCT

DT 15-MAR

DE Unknown

DE Zea ma;

OC Eukary;

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RA TOUZET

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     RESULT 38
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Best Loc
Matches
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Touzet P., Riccardi F., Morin C., Damer Touzet P., Riccardi F., Morin C., Damer Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein genome analysis program.";
Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANEOUS: On the 2D-gel the de protein is: 6.1, its MW is: 50.3 kD
                                                                                                                                                                                                                                                             MaizeDB; 123950; -.

NON_TER 1

NON_TER 14 1

SEQUENCE 14 AA; 14
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28-FEB-2003
Neurotensin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P80624;
01-OCT-1996 (
01-OCT-1996 (
15-MAR-2004 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; Liliopsida; Poal
PACCAD clade; Panicoideae; Andropogoneae; Zea.
NCBI_TaxID=4577;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -!- FUNCTION: Smooth muscle-contracting peptide.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the neurotensin famil PIR; A53608; A53608.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "[Ser7]neurotensin: isolation from guinea pig intestine."; FEBS Lett. 202:187-192(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=86248085; PubMed=3087775; Shaw C., Thim L., Conlon J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                               Maize-2DPAGE; P80624; COLEOPTILE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Small intestine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zea mays (Maize)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vasoactive; Pyrrolidone carboxylic acid.
MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID
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  (Rel. 43, Last annotation update)
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Pred. No.
1; Mismatc
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Pred. No. 9.7e+03
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Hystricognathi; Caviidae; Cavia.
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Best Local :
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"Purification and partial characterization of from rainbow trout, Oncorhynchus mykiss.";
Submitted (DEC-1999) to Swiss-Prot.
-!- FUNCTION: Antibacterial activity against G-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUB SPECIFICITY: Plasma serum.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
NCBI TaxID=8022;
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16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
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"Purification and partial characterization from rainbow trout, Oncorhynchus mykiss.";
Submitted (DEC-1999) to Swiss-Prot.
-!- FUNCTION: Antibacterial activity agains-
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Plasma serum.
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16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence up
16-OCT-2001 (Rel. 40, Last annotation
Salmocidin 2A (Fragment).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleost
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
                                                                                                                                                                                                                                                                                       Antibiotic.
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Pred. No. 1.1e
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TISSUE=Larva;
MEDLINE=98054539; PubMed=93928zz;
MEDLINE=98054539; PubMed=93928zz;
Duve H., Johnsen A.H., Maestro J.-L.,
Duvey M., East P.D., Thorpe A.;
Pavey M., East P.D., Thorpe A.;
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MOD_RES 8 8
SEQUENCE 8 AA; 898 MW.
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Cydia pomonella (Codling moth).
Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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01-OCT-1989
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   NON TER
SEQUENCE
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Peptides 18:1301-1309(1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus type 1 (Z-84 isolate) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                        EMBL; J03653; AAA44687.1; -. HIV; J03653; NEF$JY1. AIDS; Myristate; GTP-binding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=88281278; PubMed=3395517;

Josephs S.F., Reitz M.S. Jr.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Viruses; Retroid
NCBI_TaxID=11681;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'Nucleotide sequence analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       solate of HIV-
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                                                                                                                                                                                                                                                                                                                                                                               ACTIVITIES. IT MISCELLANEOUS:
                                                                                                                                                                                                                                                                                                                                                            ZAIREAN MALE.
                                                                   Myristate;
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                                                                                                                                                                                                                                                                                                                                                                               S. Hum. Retroviruses 4:165-173(1988).
ICTION: NEF has GTPase, GTP-binding and ivities. It seems to down-regulate the CELLANEOUS: THE Z-84 ISOLATE WAS TAKEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 2; Conserv
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989 (Rel. 12, Last sequence update)
983 (Rel. 42, Last annotation update)
factor (F-protein) (27 kDa protein)
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     319CB325A3733878 CRC64;
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Pred. No.
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RESULT 42
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MEDLINE=93083101; PubMed=1360367;
MEDLINE=93083101; PubMed=1360367;
Fonagy A., Schoofs L., Proost P., van Damme
"Isolation and primary structure of two sulf
the fleshfly, Neobellieria bullata.";
Comp. Biochem. Physiol. 103C:135-142(1992).
-!- FUNCTION: Myotropic peptide.
-!- SIMILARITY: Belongs to the gastrin/chole
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT; 9 AA P34966; PRT; 9 AA P34966; O1-FEB-1994 (Rel. 28, Created) O1-FEB-1994 (Rel. 28, Last sequence update) 28-FEB-2003 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SARBU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=90341105; PubMed=2381877;
Murphy R., Turner C.A., Furness J.B.,
"Isolation and microsequence analysis
from guinea pig small intestine.";
Peptides 11:613-617(1990);
"The process of the control o
                                                                                                                                                                                                                                                                                                                      NSKI SARBU STANDARD; PRT; 9 AA. P41492; 01-NOV-1995 (Rel. 32, Created) 01-NOV-1995 (Rel. 32, Last sequence update) 01-FEB-1996 (Rel. 33, Last annotation update) Neosulfakinin-I (NEB-SK-I).
                                                                                                                                                                                                                                     Sarcophaga bullata (Grey flesh fly) (Neobellie Eukaryota; Metazoa; Arthropoda; Hexapoda; Inse Neoptera; Endopterygota; Diptera; Brachycera; Sarcophagidae; Sarcophaga.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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-:- SUBCELLULAR LOCATION: Secreted.
-:- SIMILARITY: Belongs to the NmU famil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Small intestine;
                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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PROSITE; PS00967; NMU; 1.
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Pred. No.
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Hystricognathi; Caviidae; Cavia.
        gastrin/cholecystokinin family
                                                                                        van Damme
of two sulf
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1.4e+05;
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form of
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RESULT 45
HTF_HELZE
ID HTF_HELZE STANDARD; PRT; 10 AA.

AC P16353;
DT 01-AUG-1990 (Rel. 15, Created)
DT 01-FEB-1994 (Rel. 28, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Hypertrehalosaemic hormone (HeZ-HRTH).
OS Heliothis zea (Corn earworm) (Bollworm).
OC Eukaryota; Metazoa; Arthropoda; Hexapoda; Insect
OC Neoptera; Endopterygota; Lepidoptera; Glossata;
OC Noctuidae; Heliothinae; Helicoverpa.

RN [1]
RP SEQUENCE.
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Best Local S
Matches 2
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PROSITE; PS00259; GASTRIN; 1.
Neuropeptide; Amidation; Sulfat
MOD_RES 4 4 SU
MOD_RES 9 9 AM
SEQUENCE 9 AA; 1187 MW; 8B0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the type-B carboxylesterase/lipase family.
InterPro; IPR002018; CarbesteraseB.
PROSITE; PS00122; CARBOXYLESTERASE B 1; PARTIAL.
PROSITE; PS00941; CARBOXYLESTERASE B 2; PARTIAL.
Hydrolase; Serine esterase.
NON TER 10 10
SEQUENCE 10 AA; 1025 MW; 018ABE587865A2CO CRC64
                                                      Heliothis zea (Corn earworm) (Bollworm).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia; Noctuoidea;
Noctuidae; Heliothinae; Helicoverpa.
NCBI TaxID=7113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=97468499; PubMed=9327586; Siegfried B.D., Ono M., Swanson J.J.; Purification and characterization of a carboxylesterase associated with organophosphate resistance in the greenbug, Schizaphis graminum (Homoptera: Aphididae)."; Arch. Insect Biochem. Physiol. 36:229-240(1997).

-I- CATALYTIC ACTIVITY: A carboxylic ester + H(2)O = an alcohol + a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P81012;
01-NOV-1997
01-NOV-1997
15-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schizaphis graminum (Aphid).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Paraneoptera; Hemiptera; Sternorrhyncha;
Aphidoidea; Aphididae; Aphidini; Schizaphis.
NCBI TaxID=13262;
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MEDLINE=97468499;
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01-NOV-1997 (Rel. 35, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
Esterase 52 kDa subunit (EC 3.1.1.1) (Carboxylic-ester hydrolase)
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Pred. No. 1.1e+04;
0; Mismatches 0;
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AMIDATION (POTENTIAL).
BECA0691E86B5AAA CRC64;
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Best Local S
Matches 2
                                                             TISSUE=Corpora cardiaca;
MEDLINE=88326324; PubMed=3415690;
Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Bird T.G.,
Tseng C.M., Zhang Y.S., Hayes D.K.;
"Isolation and primary structure of a neuropeptide hormone
"Isolation and primary structure and adipokinetic activ
Biochem. Biophys. Res. Commun. 155:344-350(1988).
7
                1 GFG 3
                               . Similarity 2; Conserv
                                Conservative
                                        25.5%;
66.7%;
                                Score 13; DB 1; Le
Pred. No. 1.1e+04;
1; Mismatches 0;
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Search completed: August 30, 2004, 10:50:20 Job time: 3.55068 secs

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  pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 14
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2: pir2:*
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51
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| ALIGNMENTS | CHIGIS | 837141 | PH1586 | I51430 | PT0304 | PT0263 | 863492 | 847395 | S47394 | S47391 | S26548 | JTJG0 | PT0243 | C45474 | A42266 | C59137 |
| CÓ. | T H CHAIN A-D-O I | | chain V-D- | hemoglobin beta ch | Ig heavy chain CRD | | | T-cell antigen rec | , | T-cell antigen rec | T-cell receptor be | ٠, | Ig heavy chain CRD | thrombospondin 2 - | peptidylglycine mo | protein pf3 - gold |

RESULT 2
\$36678
dodecenoyl-CoA Delta-isomerase (EC 5.3.3.8) / enoyl-CoA hydratase (EC 4.2.1.17) / 3-hydr
C;Species: Mus musculus (house mouse)
C;Species: Musculus (house mouse)
C;Pecies: Musculus (house mouse)
C R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0331 Ig heavy chain CRD3 region (clone J2-121) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0331 RESULT PT0331 A;Experimental source: liver C;Function: A;Pathway: fatty acid beta-oxidation C;Keywords: carbon-oxygen lyase; fatty A; Molecule type: DNA
A; Residues: 1-13 < YAM>
A; Experimental source: B lymphocyte
C; Keywords: heterotetramer; immunoglobulin 片 밁 Ś Š Query Match Best Local S Matches 4 Query Match Best Local Matches Local Similarity nes 4; Conserv . เง σ W \vdash Similarity GYKNSKF 9 GYNNPPF 12 FGYKNSKF FGFKMGPF Conservative Conservative 45.1%; 57.1%; 45.1%; 50.0%; Score 23; DB Pred. No. 4.3e o •• Score 23; DB 2; Pred. No. 3.9e+02; Mismatches acid beta-oxidation; hydro-lyase; intramolecular DB 2; JB 2, 1.3e+02; 3; Length 14; Length 13; Indels 0 0 Gaps Gaps 0 0

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RESULT 4
PH0787

T-cell receptor alpha chain (F8) - mouse (fragment)

C;Species: Mus musculus (house mouse)

C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997

C;Accession: PH0787

R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.

J. Exp. Med. 174, 1371-1383, 1991

A;Title: T cell receptor genes in a series of class I major histocompatibility (allelic exclusion and antigen-specific repertoire.

allelic exclusion and antigen-specific repertoire.

Toferance number: PH0746; MUID:92078846; PMID:1836010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-13 < CAS>
A; Cross-references: EMBI
A; Experimental source: T
C; Keywords: T-cell recep
                                                                                                                                              C;Accession: PC4373
R;Sarig, G.; Weisman-Shomer, P.; Fry, M.
Blochem. Biophys. Res. Commun. 237, 617-623, 1997
A;Title: Telomeric and tetraplex DNA binding properties
A;Reference number: PC4371; MUID:97445086; PMID:9299414
A;Accession: PC4373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Mus musculus (house mouse)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 11-Apr-1995
C;Accession: PH1454
C;Casanova, J.L.; Martinon, F.; Gournier, H.; Barra, C.; Pannetier, C.; Regnault, A.; Kc J. Exp. Med. 177, 811-820, 1993
A;Title: T cell receptor selection by and recognition of two class I major histocompatik A;Reference number: PH1430; MUID:93171821; PMID:8436911
A;Accession: PH1454
A;Molecule type: mRNA
A;Experimental source: cytolytic T-lymphocyte
C;Superfamily: immunoglobulin homology
C;Keywords: receptor; T-cell
                                                                         A;Molecule type: protein
A;Residues: 1-8 <SAR>
C;Comment: This protein binds either
F;1-4/Domain: RNP1 #status predicted
                                                                                                                                                                                                                                                    telomeric and tetraplex DNA binding protein qTBP42 III - C;Species: Rattus norvegicus (Norway rat) C;Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_C;Accession: PC4373
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Experimental source: T ly Reywords: T-cell receptor
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                                     Query Match
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                  41.2%;
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Score 21; DB 2; Li
Pred. No. 2.8e+05;
1; Mismatches 0;
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RESULT 8
PT0211
                                                    A; Molecule type: mRNA
A; Residues: 1-11 <NAK>
C; Keywords: T-cell receptor
                                                                                                        R;Nakano, N.; Kikutani, H.; Nishimoto, H.; Kishimoto, T. J. Exp. Med. 173, 1091-1097, 1991
A;Title: T cell receptor V gene usage of islet beta cell-reactive A;Reference number: PT0209; MUID:91217621; PMID:1902501
A;Accession: PT0211
                                                                                                                                                                                                     T-cell receptor alpha chain V-J region (4-1-G.4) - C; Species: Mus musculus (house mouse) C; Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 C; Accession: PT0211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Accession: B33995
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Nac Proc. Natl. Acad. Sci. U.S.A. 86, 8161-8164, 1989
A;Title: Primary structure of two neuropeptide hormones A;Reference number: A33995; MUID:90046758; PMID:2813385 A;Accession: B33995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide;
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status predicted
F;10/Modified site: amidated carboxyl end (Tyr) #status predicted
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A; Residues: 1-9 < OCH >
C; Keywords: protein biosynthesis;
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C;Species: Tabanus atratus (black horse fly)
C;Date: 23-Mar-1990 #sequence_revision 23-Mar-1990 #text_change 31-Oct-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to JIPID, February 1994
A;Description: Phylogenetic diversity
A;Reference number: JP0042
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C;Species: Leuconostoc mesenteroides
C;Date: 10-Mar-1994 #sequence_revision 28-Oct-1994 #text_change 28-Oct-1994
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R;Ochi, K.
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C;Date: 19-May - C;Accession: S38307
C;Accession: S38307
R;Wang, G.L.; Goldstein, E.S.
R;Wang, G.L.; Goldstein, E.S.
R;Wang, G.L.; Goldstein, E.S.
R;Wang, G.L.; Goldstein, E.S.
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A; Title: Isolation of four novel tachykinins from frog (Ra: A; Reference number: JE0426; MUID:91254337; PMID:2043143
A; Accession: JE0427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R; Kangawa, K.; Kozawa, H.; Hino, J.; Minamino, N.; Matsuo, Regul. Pept. 42(Suppl.1), S12, 1992
A; Title: Isolation of four novel tachykinins from frog (Ran A; Reference number: A61033
A; Accession: B61033
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A;Title: An AP-1 binding site in the upstream region of A;Reference number: S38307; MUID:94032494; PMID:8218421
A;Accession: S38307
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                                                        36K kidney stone protein - unidentified bacterium (fragment) C;Species: unidentified bacterium C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_chacC;Accession: B58502
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R;Binette, J.P.; Binette, M.B. submitted to the Protein Sequence Database, October 1996 A;Description: The proteins of kidney and gallbladder stones
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C;Species: Tripneustes gratilla
C;Date: 17-Apr-1993 #sequence revision 17-Apr-1993 #text_change 18-Aug-2000
C;Accession: F60527; H39572
R;Yoshino, K.I.; Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita Comp. Biochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and it otus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Reference number: A60527
A;Accession: F60527
A;Accession: F60527
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R;Tsugita, A.; Kawakami, T.; Uchida, T.; Sakai, T.; Kamo, M.; Matsui, T.; Watanabe, Electrophoresis 21, 1853-1871, 2000
A;Title: Proteome analysis of mouse brain: Two-dimensional electrophoresis profiles A;Reference number: PC7072
A;Accession: PC7074
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A;Molecule type: protein
A;Residues: 1-14 <BIN>
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                                                                                                                                                                                                              A; Note: this peptide did not contain bromophenylalanine R; Yoshino, K.; Takao, T.; Suhara, M.; Kitai, T.; Hori, H.; Nomura, K.; Naiochemistry 30, 6203-6209, 1991
A; Title: Identification of a novel amino acid, o-bromo-L-phenylalanine, A; Reference number: A39572; MUID:91283461; PMID:2059627
A; Accession: H39572
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A;Experimental source: strain C57BL/6Cr
C;Keywords: brain; cerebral cortex
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                                                                                     C; Keywords: bromine F; 2/Modified site:
                                                                                                                                                                     A; Molecule type: protein A; Residues: 1-10 < YO2>
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                                                                                                                            A; Note: this peptide contained bromophenylalanine C; Superfamily: unassigned animal peptides
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;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000
;Accession: PC7074
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35.3%; Score 18; DB 2; Lilarity 100.0%; Pred. No. 2.7e+03; Conservative 0; Mismatches 0;
                                                                                   2'-bromophenylalanine (Phe) (partial) #status
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Pred. No.
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Pred. No.
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3. 2.8e+05;
0;
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3. 2.4e+03;
0;
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                                              Length 10;
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A; Molecule type: protein
A; Residues: 1-10 <OBE>
A; Experimental source: leaf,
C; Keywords: alternation
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C;Accession
R;Yoshino,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 15
$06964

hypothetical protein (nifA 5' region) - Rhizobium leguminosarum (fragment)
C;Species: Rhizobium leguminosarum
C;Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 08-Oct-1999
C;Accession: $06964
R;Roelvink, P.W.; Hontelez, J.G.J.; van Kammen, A.; van den Bos, R.C.
Mol. Microbiol. 3, 1441-1447, 1989
A;Title: Nucleotide sequence of the regulatory nifA gene of Rhizobium leguminosarum A;Accession: $06964; MUID:90136072; PMID:2693897
A;Accession: $06964
                                                                                                                                                                                     R;Obenland, D.M.; Simmen, U.; Boller, T.; Wiemken, A. Plant Physiol. 101, 1331-1339, 1993
A;Title: Purification and characterization of three soluble A;Reference number: PQ0752; MUID:94143483; PMID:8310063
                                                                                                                                                                                                                                                                                beta-fructofuranosidase (EC 3.2.1.26) IIB - barley (fragment)
C; Species: Hordeum vulgare (barley)
C; Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change
C; Accession: PQ0753
                                                                                                                                                                       A; Accession: PQ0753
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B60589
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A; Residues: 1-10 < YOS>
C; Superfamily: unassigned animal
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A;Title: A halogenated amino acid-containing sperm activating peptide and its related otus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Reference number: A60527
A;Accession: B60589
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C;Species: Heterocentrotus mamillatus
C;Species: Heterocentrotus mamillatus
C;Sate: I7-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
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Best Local S
Matches 3
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;Residues: 1-10 <ROE>
;Cross-references: EMBL:X17073; NID:g46208; PIDN:CAA34923.1;
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Best Local S
Matches 3
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3; Conserv
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3; Conserv
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ilarity 100.0%; Pred. No. 2.7e+03;
Conservative 0; Mismatches 0;
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initiators; glycoprotein; glycosidase; hydrolase
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100.0%; Pr
           35.3%;
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0; Mismatches
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           Score 18;
Pred. No.
           DB 2;
2.7e+03;
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                               Length 10;
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Ig heavy chain CRD3 region (clone 3-109B) - human (fragment) C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_char C;Accession: PT0274
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0274
A;Molecule type: DNA
A;Residues: 1-12 <YAM>
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C;Accession: S26547
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, F. J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: S26547
                                                                                                                                                                                                                                                                                                                                C;Species:
C;Date: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: protein
A;Residues: 1-11 <KRA>
C;Comment: This enzyme was purified to homogeneity and shown to have 20alpha hydroxysterc sphate dehydrogenase (GAPDH) activity although the fragment shows near identify to known C;Keywords: homotetramer; NAD; NADP; oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Krafft, A.E.; Hylemon, P.B.

J. Bacteriol. 171, 2925-2932, 1989

A;Title: Purification and characterization of a novel form A;Reference number: A44755; MUID:89255043; PMID:2722736

A;Recession: A44755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20alpha-hydroxysteroid dehydrogenase (EC 1.1.1.149) - C;Species: Clostridium scindens C;Date: 12-Mar-1993 #sequence_revision 12-Mar-1993 #te C;Accession: A44755
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A44755
                                                                                                                                                                                A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
                                 Query Match
Best Local S
Matches 3
                                                                                                                    Cross-references: EMBL:X67997
Experimental source: cytolytic T-lymphocyte, clone Cw3/10.
Superfamily: immunoglobulin V region; immunoglobulin homol
                                                                                                  Superfamily: immunogropur:
                                                                                                                                                                                                                                                                                                           -cell receptor beta chain (clone Cw3/10.1) - mouse
;Species: Mus musculus (house mouse)
;Date: 13-Jan-1995 #sequence_revision 17-Apr-1998
;Accession: S26547
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Best Local (
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75.0%;
                               Score 18; DB
Pred. No. 3.2e
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Pred. No.
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2.9e+03;
                                                                  DB 2;
                              .2e+03;
                                                                                                                                                                                                                                                                                                                                                               mouse (fragment)
                                                                                                                                                                                                                                                                                        Necker, A.; Gournier, H.; Barra,
                                                                                                                                                                                                                                                                                                                           #text_change 17-Mar-1999
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#text_change

16-Aug-1996

S.; Caton, A.J.; Rovera,

heavy

chain diversity and

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A; Experimental source: B ly C; Keywords: heterotetramer;
histone H4-1 precursor - African clawed frog (fragment)
C;Species: Xenopus laevis (African clawed frog)
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change
C;Accession: I51432
R;Gargiulo, G.; Razvi, F.; Ruberti, I.; Mohr, I.; Worcel, A.
J. Mol. Biol. 181, 333-349, 1985
A;Title: Chromatin-specific hypersensitive sites are assembled o
A;Reference number: I51431; MUID:85160855; PMID:4039007
A;Accession: I51432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        glyceraldehyde-3-phosphate dehydrogenase (phosphorylating) (EC 1.2.1.12) - rice C;Species: Oryza sativa (rice) C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 03-Jun-2002 C;Accession: PS0277
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A,Note: VCAM-1 5' untranslated region binding protein with a probable translation
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A;Accession: A59387
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                                                                                                                                                                                                                                                                                                                                                                                                                                         submitted to JIPID,
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Best Local S
Matches 3
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Best Local (
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Best Local (
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3; Conserva
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Pred. No. 3.5e+03;
0; Mismatches 0;
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Pred. No. 3.2e+03;
1; Mismatches 1
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of a novel VCAM-1 5'UTR
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A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-14 <GAR>
A;Cross-references: GB:M23777; NID:g214219; PIDN:AAA49737.1; PID:g214222
C;Superfamily: histone H4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             glucose 1-dehydrogenase (NAD) (EC 1.1.1.118) - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 03-Jun-2002
                                                                                                                                                                                                                                                                                                                                                                                               R;Charlier, D.; Piette, J.; Glansdorff, N.
Nucleic Acids Res. 10, 5935-5948, 1982
A;Title: is3 can function as a mobile promoter in e.coli.
A;Reference number: I41137; MUID:83064529; PMID:6292860
A;Accession: I41138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type:
A; Residues: 1-14
C; Keywords: NAD;
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A;Title: Amino acid sequence of the tryptic
A;Reference number: A17150; MUID:82182061; F
A;Accession: A17150
spermadhesin AQN-3 - pig (fragments)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 04-Dec-1997 #sequence_revision 12-Dec-1997 #
C;Accession: S68649
R;Calvete, J.J.; Dostalova, Z.; Sanz, L.; Adermann,
FEBS Lett. 379, 207-211, 1996
A;Title: Mapping the heparin-binding domain of boar
                                                                                                                       RESULT 25
S68649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Escherichia coli
C;Date: 31-May-1996 #sequence_revision 31-May-1996 #text_change 08-Oct-1999
C;Accession: I41138
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                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-11 < RES>
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spermadhesins

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A; Reference number: S68648; A; Accession: S68649 A; Status: preliminary A; Molecule type: protein A; Residues: 1-6;7-11 < CAL>
Ig H chain V-D-J region (wild-type clone 304) - mon C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994
C;Accession: PH1597
R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
                                                                                                                                                   RESULT 28
PH1597
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PH0939
T-cell receptor beta chain V-D-J region (clone 10) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
J. Experimental norvegicus (Norway rat)
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental A;Fitter norvegicus (Norway rat)
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental A;Fitter norvegicus (Norvegicus (Norvegic
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peptidylglycine monooxygenase (EC 1.14.17.3), rPAM-N;Alternate names: peptidylglycine alpha-amidating C;Species: Rattus norvegicus (Norway rat) C;Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 C;Accession: A42266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unidentified 27.2K protein - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 18-Aug-2000
C;Accession: PC7079
R;Tsugita, A.; Kawakami, T.; Uchida, T.; Sakai, T.; Kamo, M.; Matsui, T.; Watanabe, Electrophoresis 21, 1853-1871, 2000
A;Title: Proteome analysis of mouse brain: Two-dimensional electrophoresis profiles
A;Reference number: PC7072
                                                                                                                                                                                                                                                                                                                                                                                              R;Sakamoto, Y.; Ando, A.; Tamai, Y.; Miura, K. submitted to the Protein Sequence Database, November 1999 A;Description: Differences of proteins expressed in the fi
                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Accession: C59137
R;Sakamoto, Y.; Ando,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein Pf3 - golden needle mushroom (fragment)
C;Species: Flammulina velutipes (golden needle mushroom)
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text
                                                                                              A42266
                                                                                                                                                                                                                                                                                                                         A; Molecule type: protein A; Residues: 1-14 < SAK>
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A; Accession: C59137
                                                                                                               RESULT 31
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A;Title: Molecular characterization of transgene-induced immunodeficiency A;Reference number: PH1580; MUID:93301609; PMID:8315387
A;Accession: PH1597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Experimental source: strain C57BL/6Cr Slc,
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A;Residues: 1-14 <TSU>
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C; Keywords: immunoglobulin
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A; Residues: 1-14 <LEV>
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Matches 3
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Best Local !
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3; Conservative
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1; Mismatches
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Pred. No. 5.7e+03;
0; Mismatches :
               18-Nov-1994 #text_change 26-May-1995
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5.7e+03;
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RESULT 32
C45474
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A;Title: Alternative splicing and endoproteolytic processing A;Reference number: A42266; MUID:92156145; PMID:1740449
A;Accession: A42266
A;Status: preliminary
                                                                                                                                                                                                                                                R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain div. A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0243
                                                                                                                                                                                                                                                                                                                            Ig heavy chain CRD3 region (clone 2-103A) - human (fragment) C; Species: Homo sapiens (man) C; Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_cha C; Accession: PT0243 R; Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Cato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N;Alternate names: corticotropin-induced secreted protein (CISP); thrombospondin C;Species: Bos primigenius taurus (cattle) C;Date: 21-Sep-1993 #sequence_revision 18-Nov-1994 #text_change 14-Aug-1998 C;Accession: C45474
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A;Note: sequence extracted from NCBI backbone (NCBIN:82733, NCBIP:82750)
C;Keywords: oxidoreductase
                                                                                                                                                                     A;Molecule type: DNA
A;Residues: 1-10 <YAM>
A;Experimental source: B lyr
C;Keywords: heterotetramer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: protein A; Residues: 1-10 < PEL>
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A;Title: Corticotropin-induced secreted protein, an ACTH-induced protein secreted by A;Reference number: A45474; MUID:93179438; PMID:8382699
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Pred. No. 6.3e+03;
1; Mismatches 0
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Pred. No. 2.8e+05;
3; Mismatches (
                                                                                          Score 16; DB:
Pred. No. 6.3e
1; Mismatches
                                                                                                                                                                                                                                                                        fic immunoglobulin heavy chain diversity PMID:1899102
                                                                                                              .3e+03;
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A;Description:
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A; Molecule type: protein
A; Residues: 1-12 < SAK1 >
A; Note: the farnesyl may instead be (2E, 6E
R; Sakagami, Y.; Isogai, A.; Suzuki, A.; Ta
Agric. Biol. Chem. 42, 1301-1302, 1978
A; Title: Amino acid sequence of tremerogen
A; Reference number: A61313
A; Accession: A61313
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R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific A;Reference number: S26512; MUID:92364546; PMID:1380061 A;Accession: S26548
C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995
C;Accession: S47391; S47386
R;Lehner, P.J.
submitted to the EMBL Data Library, August 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Superfamily: tremerogen a-13
C;Keywords: extracellular protein; hormone; lipoprotein; methylated carboxyl end; pherom: F;12/Binding site: farnesyl (Cys) (covalent) #status experimental F;12/Modified site: methyl ester carboxyl end (Cys) #status experimental
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Science 212, 1525-1527, 1981
A;Title: Peptide sex hormones inducing conjugation tube formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tremerogen A-10 - jelly fungus (Tremella mesenterica)
C;Species: Tremella mesenterica
C;Date: 03-Aug-1984 #sequence revision 03-Aug-1984 #text_change 07-Nov-1997
C;Accession: A01642; A61313
                                                                                                                                      RESULT
S47391
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A;Experimental source: cytolytic T-lymphocyte, clone Cw3/Cas3 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T-cell receptor beta chain (clone Cw3/Cas3) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 13-Jan-1995 #sequence_revision 17-Apr-1998 #text_change 17-Mar-1999
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A; Residues: 1-12 < CAS>
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A;Accession: A01642
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                                                                                                                T-cell antigen receptor VJ junction beta chain - human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tremerogen A-10,
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Pred. No. 7.6e+03;
0; Mismatches 1
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                        #text_change
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R;Lehner, P.J.
submitted to the EMBL Data Library, August
 RESULT 39
$63492
dissimilatory sulfite reductase beta chain, C;Species: Desulfovibrio desulfuricans
                                                                                                                                                                                                                                                                                                                          T-cell antigen receptor VJ junction beta chain - human C;Species: Homo sapiens (man) C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change C;Accession: S47395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-12 <LEH>
A;Cross-references: EMBL:Z35712;
C;Keywords: T-cell receptor
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                                                                                                                                                                                                                                                              A;Reference number: S47355
A;Accession: S47395
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A;Description: Human HLA-A0201 restricted recog
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number: S47355
A; Accession: S47394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T-cell antigen receptor VJ junction beta chain - human C;Species: Homo sapiens (man) C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change C;Accession: S47394; S47369
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A; Accession: $47391
A; Status: prelimina:
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Matches 2
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Matches 2
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                                                                                                                                                                                        ;Cross-references: EMBL:Z35715; NID:g527525; PIDN:CAA84784.1; PID:g527526;Keywords: T-cell receptor
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larity 66.7%;
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                                                                                                                                      Score 16; DB 2;
Pred. No. 7.6e+03;
1; Mismatches 0
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Pred. No. 7.6e+03;
1; Mismatches (
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Pred. No. 7.6e+03;
1; Mismatches C
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              RESULT 42
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beta

chain

African

clawed

frog

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Ig heavy chain CRD3 region (clone 5-115B) - human (fragment) C;Species: Homo sapiens (man) C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_cha C;Accession: PT0304 R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Cato J. Exp. Med. 173, 395-407, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0263
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0263
A;Accession: PT0263
A;Residues: 1-13 <YAM>
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PT0263
                                                                           A; Molecule type: DNA
A; Residues: 1-13 < YAM>
A; Experimental source: B lymphocyte
C; Keywords: heterotetramer; immunoglobulin
                                                                                                                                                        R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Recession: PT0304
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A;Molecule type: protein
A;Residues: 1-13 <STE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999 C;Accession: S63492
R;Steuber, J.; Arendsen, A.F.; Hagen, W.R.; Kroneck, P.M.H.
Eur. J. Biochem. 233, 873-879, 1995
                                                                                                                                                                                                                                                                                                                                                                    RESULT 41
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Query Match
Best Local S
Matches 2
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Keywords: heterotetramer;
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er; immunoglobulin
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75.0%;
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Score 16; DB 2; Louis Pred. No. 8.2e+03; 1; Mismatches 0;
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                                    Length 13;
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C;Accession: I51430
R;Kay, R.M.; Harris, R.; Patient, R.K.; Williams, J.G.
Nucleic Acids Res. 8, 2691-2707, 1980
A;Title: Molecular cloning of cDNA sequences coding for A;Reference number: I51428; MUID:81053706; PMID:7001356
A;Accession: I51430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice A;Reference number: PH1580; MUID:93301609; PMID:8315387
A;Accession: PH1580
                                                                                                                                                                                                     R;Douillie, A.; Toussaint, A.; Faelen, M. submitted to the EMBL Data Library, August 1993 A;Description: Identification of the integration host factor genes A;Reference number: S37139 A;Accession: S37141
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A;Residues: 1-14 <LEV>
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A;Residues: 1-14 <KAY>
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                                                                                                                          A;Cross-references: EMBL:X74750; NID:g399669; PIDN:CAA52769.1;
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                                          Query Match
Best Local S
Matches 2
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                                                                                                                                                                                                                                                                                                            Species: Erwinia chrysanthemi;Date: 06-Jan-1995 #sequence_revision;Accession: S37141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      H Chain V-D-J region (wild-type clone 10) - mouse (fragment)
Species: Mus musculus (house mouse)
Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
Accession: PH1586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Experimental source: bone marrow pre-B lymphocyte Keywords: immunoglobulin
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     4 YKNSK 8
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Similarity 66.7%;
2; Conservative
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Pred. No. 8.8e+03;
1; Mismatches
                                                               Score 15; DB 2;
Pred. No. 2.8e+05;
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                                                                                                                                                                                                                                                                                                                                    06-Jan-1995 #text_change 08-Oct-1999
                                                                                    Length 8;
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FKSAK 6

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PH1618
Ig H Chain V-D-7 region (clone B-less 33) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
C;Accession: PH1618
R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
J. Exp. Med. 178, 317-329, 1993
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice A;Reference number: PH1580; MUID:93301609; PMID:8315387
A;Accession: PH1618
A;Molecule type: DNA
A;Residues: 1-8 <LEV>
A;Experimental source: bone marrow pre-B lymphocyte
C;Keywords: immunoglobulin
Query Match
Best Local Similarity 66.7%; Pred. No. 2.8e+05;
Matches 2; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
Matches 2; Conservative 1; Mismatches 0; Indels 0; Gaps 5
GYG 7
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Search completed: August 30, 2004, 10:58:48

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Perfect score:
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| length: 14
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                   August 30, 2004, 10:49:45; Search time 11.0068 Seconds (without alignments) 257.253 Million cell updates/sec
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      Published_Applications_AA:*
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: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
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  14 US-10-239-313A-642
14 US-10-239-313A-542
14 US-10-239-313A-680
B US-08-996-140-15
US-09-876-904A-78
110 US-09-876-904A-78
1110 US-09-26-795-15
112 US-09-963-658-263
113 US-09-963-693-263
114 US-09-963-693-263
115 US-09-963-693-263
116 US-09-983-290-313
117 US-10-449-829A-28
118-09-988-493-274
119 US-10-014-340-467
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US-09-226-666-4
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Sequence 13, Appli
Sequence 4, Appli
Sequence 642, App
Sequence 542, App
Sequence 580, Appl
Sequence 15, Appl
Sequence 78, Appl
Sequence 263, App
Sequence 274, App
Sequence 313, App
Sequence 274, App
Sequence 274, App
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| -09-799-250-33 | -09-799-250-23 | -09-799-250-15 | -10-351-641-9 | 0-174-105A-1 | -10-474-601-1 | -10-302-100B- | -10-317-550- | -09-572-270A- | -10-807-856-6 | -10-003-983B | -10-460-512-2 | -10-460-512- | 09-879-957-6 | -10-387-486-1 | -10-014-485A- | -10-013-815- | -10-013-815-3 | -10-319-340 | -09-780-053-47 | 09-780-053-3 | 09-780-053-30 | 09-780-053-18 | -10-154-884B-1094 | -10-057-475B- | 9-780-053-158 | S-10-264-309 | -10-255-679-6 | -09-791-377-52 | -10-014-340-4 | _ |
| equence 33 | equence 23 | equence 15 | equence 91 | quence 15 | equence 12 | equence 28 | equence 2, | equence 62 | equence 63 | equence 11 | equence 28 | equence 6, | quence 63, | equence 10 | equence 11 | e 38 | equence 35 | equence 8, | equence 472 | e 399 | equence 304 | e H | equence 10 | equence 1094 | quence 158, | Sequence 24 | equence 60, | e 526, | equence 468, | |

ALIGNMENTS

RESULT 1 US-10-421-503-13

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RESULT 4
US-10-239-313A-542
; Sequence 542, Application US/10239313A
; Publication No. US20030175285A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Clostridium tetani
US-10-239-313A-642
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-239-313A-642
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 642
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: KLINGUER - HAMOUR, Christine
APPLICANT: CORVAIA, Nathalie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 642, Application US/10239313A Publication No. US20030175285A1
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Best Local Similarity
APPLICANT: KLINGUER - HAMOUR, Christine
APPLICANT: CORVAIA, Nathalie
APPLICANT: BECK, Alain
APPLICANT: BECK, Liliane
TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN T
TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID
FILE REFERENCE: 343 727 - US
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APPLICANT: GOETSCH, Liliane
TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT
TITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE
TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID
FILE REFERENCE: 343 727 - US
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TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
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ORGANISM: Artificial Sequence
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APPLICANT: KLINGUER - HAMOUR, Christine

APPLICANT: KLINGUER - HAMOUR, Christine

APPLICANT: CORVAIA, Nathalie

APPLICANT: CORVAIA, Nathalie

APPLICANT: BECK, Alain

APPLICANT: BECK, Alain

APPLICANT: GOETSCH, Liliane

TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITTLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID

TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID

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TITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID

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; Sequence 15, Application US/08996140
; Publication No. US20030190318A1
; GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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PRIOR FILING DATE: 2000-03-23
PRIOR APPLICATION NUMBER: PCT 01/70772
PRIOR FILING DATE: 2001-03-22
PRIOR FILING DATE: 2001-03-22
PRIOR SEQ ID NOS: 697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/239,313A CURRENT FILING DATE: 2002-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PR
                                                                                                APPLICANT: TORIGOE, Kakuji
APPLICANT: USHIO, Shimpei
APPLICANT: KUNIKATA, Toshio
APPLICANT: KURIMOTO, Masashi
TITLE OF INVENTION: INTERLEUKIN-18 RECEPTOR PROTEINS
NUMBER OF SEQUENCES: 31
                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AN
   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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BROWDY AND NEIMARK
419 Seventh Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49.0%; Score 25; DB 14; ilarity 100.0%; Pred. No. 1.2e+06 Conservative 0; Mismatches
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Suite 300
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hes 0;
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SEQUENCE CHARACTERISTICS:
SEQUENCE TO Amino acids
                                                                                                                                SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 78
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 78, Application US/09876904A Publication No. US20030072794A1
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                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                           APPLICANT: BOULIKAS, TENI
TITLE OF INVENTION: ENCAPSULATION OF PLASMID DNA (LIPOGENES TM) AND THERAPEUTIC
TITLE OF INVENTION: AGENTS WITH NUCLEAR LOCALIZATION SIGNAL/FUSOGENIC PEPTIDE
TITLE OF INVENTION: CONJUGATES INTO TARGETED LIPOSOME COMPLEXES
FILE REFERENCE: TB-2002.00
CURRENT APPLICATION NUMBER: US/09/876,904A
CURRENT FILING DATE: 2001-06-08
CURRENT FILING DATE: 2001-06-08
                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/210,925
PRIOR FILING DATE: 2000-06-09
NUMBER OF SEQ ID NOS: 629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: lir
MOLECULE TYPE:
FRAGMENT TYPE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide crosslinked OTHER INFORMATION: bovine serum albumin
                                                ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: BROWDY, Roger L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: TO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: JP 163,490/1997 FILING DATE: 6-JUN-1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: JP 52,526/1997
FILING DATE: 21-FEB-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/996,140 FILING DATE: 22-DEC-1997 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: JP 2: FILING DATE: 28-JUL-1997 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: JP 356,426/1996 FILING DATE: 26-DEC-1996
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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internal fragment
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25,618
TORIGOE=2
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US-09-205-658-263

(Sequence 263, Application US/09205658

; Patent No. US20010029617A1

; GENERAL INFORMATION:

APPLICANT: Ruvkun, Gary

APPLICANT: OGG, Scott

TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS

FILE REFERENCE: 00786/351004

CURRENT APPLICATION NUMBER: US/09/205,658

CURRENT FILING DATE: 1998-12-03

; EARLIER APPLICATION NUMBER: 08/857,076

EARLIER APPLICATION NUMBER: 08/88,534

; EARLIER APPLICATION NUMBER: US98/10080

: EARLIER APPLICATION NUMBER: US98/10080
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                                                                  ; ORGANISM: Caenorhabditis elegans US-09-205-658-263
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: HART, MARY KATE
APPLICANT: WILSON, JULIE
TITLE OF INVENTION: MONOCLONAL ANTIBODIES AND COMPLEMENTARITY-DETERMINING
TITLE OF INVENTION: REGIONS BINDING TO EBOLA GLYCOPROTEIN
                                                                                                 NUMBER OF SEQ ID NOS: 328
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 263
LENGTH: 13
TYPE: PRT
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Best Local :
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                Query Match
Best Local
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 Matches
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CURRENT APPLICATION NUMBER: US/10/226,795
CURRENT FILING DATE: 2002-11-18
                                                                                                                                                                                     EARLIER FILING DATE: 1998-05-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence
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Score 25; DB Pred. No. 6e+0

1; Mismatches
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Pred. No.
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Pred. No.
                DB 9;
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5.5e+02;
                                 Length 13;
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RESULT 10
US-09-963-693-263
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; Publication No.
RESULT 12
US-09-826-290-313
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US-09-791-378-526
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CURRENT FILING DATE: 2001-09-25
PRIOR APPLICATION NUMBER: US/09/205,658
PRIOR APPLICATION NUMBER: US/09/205,658
PRIOR APPLICATION NUMBER: 08/857,076
PRIOR FILING DATE: 1997-05-15
PRIOR APPLICATION NUMBER: 08/888,534
PRIOR FILING DATE: 1997-07-07
PRIOR APPLICATION NUMBER: US98/10080
PRIOR APPLICATION NUMBER: US98/10080
PRIOR APPLICATION NUMBER: US98/10080
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Best Local S
Matches 4
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 263
                                                                                                                                                                       Matches
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APPLICANT: Parekh, Rajesh
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APPLICANT: Ruvkun, Gary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 526, Application US/09791378
Patent No. US20020142303A1
                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: PROTEINS, GENES AND THBIR USE FOR DIAGNOSIS AND TREATMENT OF TITLE OF INVENTION: SCHIZOPHRENIA FILE REFERENCE: 9195-061-999 CURRENT APPLICATION NUMBER: US/09/791,378 CURRENT FILING DATE: 2001-02-23 PRIOR APPLICATION NUMBER: 09/750,395 PRIOR FILING DATE: 2000-12-28
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FITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC TOOLS FOR

FITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS

FILE REFERENCE: 00786/351004
                                                                                                                                                                                             Local Similarity
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                                                                                                                             2 FGYKON 6
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                                                                                    FGYSN 13
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                                                                                                                                                                                         49.0%;
80.0%;
                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                       Score 25; DB 9;
Pred. No. 6.4e+02;
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NUMBER OF SEQ ID NOS: 49
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 28
; LENGTH: 14
; TYPE: PRT
; ORGANISM: Homo sapiens
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Best Local Similarity
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US-10-449-829A-28
                                                                                                                                            US-10-449-829A-28
                                                          Query Match
Best Local S
Matches 4
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Publication No. US20040043425A1
GENERAL INFORMATION:
APPLICANT: TERRETT, Jonathan Alexander
APPLICANT: HALL, Ian Philip
TITLE OF INVENTION: PROTEINS, GENES AND THEIR USE FOR DIAGNOSIS AND TREATMENT OF CHROITILE OF INVENTION: ASTHMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Williams, Stephen A.
TITLE OF INVENTION: Nucleic Acid Molecules, Polypeptides and
TITLE OF INVENTION: Uses Therefor, Including Diagnosis and Treatment of
TITLE OF INVENTION: Alzheimer's Disease
FILE REFERENCE: 2572-1-001 N2
CURRENT APPLICATION NUMBER: US/09/826,290
CURRENT FILING DATE: 2001-04-30
PRIOR APPLICATION NUMBER: US 60/194,504
PRIOR APPLICATION NUMBER: US 60/194,504
PRIOR APPLICATION NUMBER: US 60/253,647
PRIOR APPLICATION NUMBER: US 60/253,647
PRIOR FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 492
NUMBER OF SEQ ID NOS: 492
                                                                                                                                                                                                                                                                               FILE REFERENCE: 2543-1-029

CURRENT APPLICATION NUMBER: US/10/449,829A

CURRENT FILING DATE: 2003-05-30

PRIOR APPLICATION NUMBER: PCT/GB01/05476

PRIOR FILING DATE: 2001-12-10

PRIOR APPLICATION NUMBER: P32060GB

PRIOR FILING DATE: 2000-12-08
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Patent No. US20020164668A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLICANT: Durham, L.Kathryn
PPLICANT: Friedman, David L
                                                            Local Similarity 80.
2 FGYKN 6
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land, P. Trey
id, Robert Reid
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B. Michael
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       49.0%;
80.0%;
                                                                            49.0%;
                                                          Score 25; DB 12;
Pred. No. 6.4e+02;
0; Mismatches 1
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Pred. No. 6.4e+02;
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                                                                                                  DB 12; Length 14;
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CURRENT APPLICATION NUMBER: US/09/988,493;
CURRENT FILING DATE: 2002-05-21
PRIOR APPLICATION NUMBER: PCT/GB01/01219
PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: GB 0006695.1
PRIOR FILING DATE: 2000-03-20
PRIOR APPLICATION NUMBER: GB 0007265.2
PRIOR APPLICATION NUMBER: GB 0007265.2
PRIOR FILING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 308
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 274
RESULT 16
US-10-014-340-468
; Sequence 468, Application US/10014340
; Publication No. US20030064411A1
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US-10-014-340-467
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                                                                                                                                                                                                                                                                   ; TYPE: PRT ; CRGANISM: Homo sapiens US-10-014-340-467
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Best Local S
Matches 4
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CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 823
SOFTWARE: PatentIn version 3.0
SEQ ID NO 467
LENGTH: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 467, Application US/10014340
Publication No. US20030064411A1
GENERAL INFORMATION:
APPLICANT: Herath, et al
TITLE OF INVENTION: Nucleic Acid Molecules, Polypeptides and Uses Therefor,
TITLE OF INVENTION: Diagnosis and Treatment of Alzheimer's Disease
FILE REFERENCE: 9195-078
                                                                                                                                                                                             Matches
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APPLICANT: O'Hare, Michael John
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TYPE: PRT
ORGANISM: homo sapien
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4; Conservative
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                                                                                                                   FGYSN 13
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Waterfield, Michael Derek
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o. US20030064419A1
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80.0%;
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Pred. No. 6.4e+02;
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Pred. No. 6.4e+02
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CURRENT APPLICATION NUMBER: US/10/014,340
CURRENT FILING DATE: 2001-12-10
NUMBER OF SEQ ID NOS: 823
SOFTWARE: PatentIn version 3.0
SEQ ID NO 468
LENGTH: 14
TYPE: PRT
ORGANISM: Homo sapiens
US-10-014-340-468
APPLICANT: BAB, ITAI
APPLICANT: BAB, ITAI
APPLICANT: GAZIT, DAN
APPLICANT: YU-CHEN, CHEN
APPLICANT: MUHLRAD, ANDRAS
APPLICANT: MUHLRAD, ANDRAS
APPLICANT: CHOREV, MICHAEL
APPLICANT: CHOREV, MICHAEL
TITLE OF INVENTION: SYNTHETIC PEPTIDES AND P
TITLE OF INVENTION: ACTIVITY AND PHARMACEUT
TITLE OF INVENTION: SAME
FILE REFERENCE: 331949-140798
CURRENT APPLICATION NUMBER: US/10/255,679
CURRENT FILING DATE: 2002-09/150,621
PRIOR APPLICATION NUMBER: US/9/150,621
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US-09-791-377-526
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Best Local S
Matches 4
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APPLICANT: Parekh, Rajesh
TITLE OF INVENTION: PROTEINS, GENES AND THEIR USE FOR DIAGNOSIS AND TREATMENT
TITLE OF INVENTION: SCHIZOPHRENIA
FILE REFERENCE: 9195-060-999
CURRENT APPLICATION NUMBER: US/09/791,377
CURRENT FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 09/750,395
PRIOR APPLICATION NUMBER: 09/750,395
PRIOR FILING DATE: 2000-12-28
NUMBER OF SEQ ID NOS: 677
SOFTWARE: PatentIn version 3.0
SEQ ID NO 526
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Best Local Similarity 80.v
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Publication No. US20030069175A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 14
TYPE: PRT
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                                                                                                       PEPTIDES AND PSEUDOPEPTIDES HAVING OSTEOGENIC AND PHARMACEUTICAL COMPOSITIONS CONTAINING THE
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Pred. No. 6.4e
0; Mismatches
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Pred. No.
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RESULT 20
US-09-780-053-158
; Sequence 158, Application US/09780053
; Patent No. US20020102640A1
; GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-10-264-309-240
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Best Local Similarity
Matches 4; Conserv
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Publication No.
                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: POA-002.01
CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT FILING DATE: 2002-10-03
PRIOR APPLICATION NUMBER: 60/326,708
PRIOR FILING DATE: 2001-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 117426
PRIOR FILING DATE: 1996-03-10
NUMBER OF SEQ ID NOS: 61
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                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 491
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               PPLICANT: TOWNSEND, ROBERT R.
PPLICANT: WHITE, W. FROST
PPLICANT: WILLIAMS, STEPHEN A.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES,
TITLE OF INVENTION: NUCLULING DIAGNOSIS AND
                                                                                                                                                                                                                                                                               ENGTH: 14
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SUNDERLAND, P. INC.
SUNDERLAND, ROBERT R.
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o. US20040022794A1
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DAVID M.
CHRISTIAN
B. MICHAEL
PETER J.
HOLLY D.
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                                                                                                                                                                                      49.0%;
80.0%;
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100.0%; Pred. No. 6.4e+02;
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                                                                                                                                                                                      Score 25; DB 16;
Pred. No. 6.4e+02;
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D TREATMENT OF ALZHEIMER'S DISEJ
                                                                                                                                                                                                   Length 14;
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Query Match
Best Local Similarity
"~+~hes 4; Conserve
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                                                                CURRENT APPLICATION NUMBER: US/10/057,475B
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: US/60/186,126
PRIOR APPLICATION NUMBER: US 60/186,126
PRIOR FILING DATE: 2000-03-01
PRIOR APPLICATION NUMBER: US 60/200,545
PRIOR APPLICATION NUMBER: US 60/200,545
PRIOR APPLICATION NUMBER: US 60/200,545
PRIOR APPLICATION NUMBER: US 60/200,303
PRIOR APPLICATION NUMBER: US 60/200,779
PRIOR APPLICATION NUMBER: US 60/200,779
PRIOR APPLICATION NUMBER: US 60/200,999
PRIOR APPLICATION NUMBER: US 60/200,999
PRIOR APPLICATION NUMBER: US 60/202,084
PRIOR APPLICATION NUMBER: US 60/202,084
PRIOR FILING DATE: 2000-05-04
PRIOR APPLICATION NUMBER: US 60/206,201
PRIOR FILING DATE: 2000-05-02
PRIOR FILING DATE: 2000-05-04
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LENGTH: 9
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APPLICANT: Mannion, Jane
APPLICANT: Clapper, Jonathan
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TITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCER
FILE REFERENCE: 129.5USU1
CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Corixa Corporation
TITLE OF INVENTION: Compositions and Methods for the Detection,
TITLE OF INVENTION: Hematological Malignancies
TILE REFERENCE: 014058-0144028-014676-7-175
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APPLICANT: Daniel E.H. Afar
APPLICANT: Pia M. Challita-Eid
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                       Prior Application data removed - SEQ ID NOS: 10979
FastSEQ for Windows Version 3.0
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Carter, Lauren
McNeill, Patricia Dianne
Corixa Corporation
and
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Steve Chappell Mitchell
Aya Jakobovits
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57.1%;
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Pred. No. 1.2e+0
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                                                       See File Wrapper or PALM.
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US-10-154-884B-10940
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                                                                                                                                                              US-10-154-884B-10940
                                                                                                                                                                                                                                                                                  Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 11290
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 10940
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Best Local Similarity 100.
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Mannion, Jane
APPLICANT: Retter, Marc W.
APPLICANT: Corixa Corporation
TITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therapy
TITLE OF INVENTION: Hematological Malignancies
FILE REFERENCE: 014058-014058-10521US
CURRENT APPLICATION NUMBER: US/10/154,884B
                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/218,950 PRIOR FILING DATE: 2000-07-14 PRIOR APPLICATION NUMBER: US 60/222,903 PRIOR FILING DATE: 2000-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/190,479
PRIOR FILING DATE: 2000-03-17
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                                                                                                                                                                                                        LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                             FEATURE:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 60/200,999
FILING DATE: 2000-05-01
APPLICATION NUMBER: US 60/202,084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 60/200,545 FILING DATE: 2000-04-27 APPLICATION NUMBER: US 60/200,303
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 2000-05-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 60/200,779
FILING DATE: 2000-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/206,201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2000-05-04
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                                                                                                  Similarity
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                                       FGYK 5
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                                                                              Conservative
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                                                                                                                                                                               Ly1484 short MHC HLA A3 class I binding peptide
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                                                                          47.1%; Score 24; DB 15;
100.0%; Pred. No. 1.2e+06;
cive 0; Mismatches 0;
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1.2e+06;
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                                                                                                                Length 9
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RESULT

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US-09-780-053-399

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TITLE OF INVENTION: 83P5G4: A TISSUE SPECIF
TITLE OF INVENTION: HIGHLY EXPRESSED IN PRO
FILE REFERENCE: 129.5USU1
CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: PRT; ORGANISM: Homo Sapiens US-09-780-053-180
片
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                                                                                Query Match
Best Local S
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SEQ ID NO 304
LENGTH: 10
TYPE: PRT
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                                                                     Matches
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Best Local Similarity
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                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
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APPLICANT: Daniel E.H. Afar
APPLICANT: Pia M. Challita-Eid
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APPLICANT: Daniel E.H. Afar
APPLICANT: Pia M. Challita-Eid
                                                                                                                                                                                                                                                                                                                                APPLICANT: Aya Jakobovits
TITLE OF INVENTION: 83P5G4: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCER
FILE REFERENCE: 129.5USU1
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1 CHONSTF 7
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D. US20020102640A1
                                  3 GYKNSKF 9
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                                                                 Similarity 4; Conserv
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Steve Chappell Mitchell
Aya Jakobovits
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Steve Chappell Mitchell
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20020102640A1
                                                                  Conservative
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                                                                   Score 24; DB Pred. No. 6.9e 2; Mismatches
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Pred. No.
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6.9e+02;
                                                                                                  Length 10;
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                                                                  0;
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TITLE OF INVENTION: 83P5G4: A TISSUB SPECIF; TITLE OF INVENTION: HIGHLY EXPRESSED IN PROFILE REFERENCE: 129.5USU1
CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 399
LENGTH: 10
; Sequence 8, Application US/10319340
                 RESULT 27
US-10-319-340-8
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US-09-780-053-472
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US-09-780-053-472
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US-09-780-053-399
                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 472
                                                                                                                                                            Matches
                                                                                                                                                                                            Query Match
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Best Local Similarity
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Patent No. US20020102640A1
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Patent No. US20020102640A1
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APPLICANT: Danie
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APPLICANT: Steve Chappell Mitchell
APPLICANT: Aya Jakobovits
FITLE OF INVENTION: 83P5G4: A TISS
FITLE OF INVENTION: HIGHLY EXPRESS
FILE REFERENCE: 129.5USU1
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PLICANT: Daniel E.H. Afar
PLICANT: Pia M. Challita-Eid
                                                                                                                                                                            Local Similarity
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                                                                                        GHQNSTF 10
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Steve Chappell Mitchell
Aya Jakobovits
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Pia M. Challita-Eid
Mary Faris
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Pred. No.
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Pred. No. 6.9e+02;
2; Mismatches 1
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APPLICANT: Heavner, George A.
APPLICANT: McEver, Rodger P.
APPLICANT: Geng, Jian-Guo
ITILE OF INVENTION: Peptide Inhibitors of Inflammation Mediated by S.
FILE REFERENCE: CTC 102 COD DIV
CURRENT APPLICATION NUMBER: US/10/319,340
CURRENT FILING DATE: 2002-12-13
PRIOR APPLICATION NUMBER: 08/135,319
PRIOR FILING DATE: 1993-10-12
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
SEQ ID NO 8
LENGTH: 10
TYPE: POT
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Sequence 35, Application US/10013815

Publication No. US20030105000A1

GENERAL INFORMATION:

APPLICANT: Pero, Stephanie
APPLICANT: Krag, David

APPLICANT: Oligino, Lyn

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITING GRB7

FILE REFERENCE: V0139/7048 (HCL/MAT)

CURRENT APPLICATION NUMBER: US/10/013,815

CURRENT FILING DATE: 2001-11-05

PRIOR APPLICATION NUMBER: US 60/245,755

PRIOR FILING DATE: 2000-11-03

NUMBER OF SEQ ID NOS: 194

SOFTWARE: PatentIn version 3.1

LENGTH: 11

TUDES: DEPTHERMENTATION OF THE PATENT 
                                                                                                                                                                                                     US-10-013-815-38
; Sequence 38, Application US/10013815
; Publication No. US20030105000A1
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Best Local Similarity
Thes 4; Conserve
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                                                                                                                                                                                                                                                                                                             RESULT 29
GENERAL INFORMATION:

APPLICANT: Pero, Stephanie
APPLICANT: .Krag, David
APPLICANT: .Oligino, Lyn
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITING GRB7
FILE REFERENCE: V0139/7048 (HCL/MAT)
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
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Pred. No. 7.6e+02;
0; Mismatches 1.
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Pred. No. 6.9e+02;
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RESULT 31
US-10-387-486-10
; Sequence 10, Application US/10387486
; Publication No. US20040001867A1
; GENERAL INFORMATION:
; APPLICANT: INSTITUT PASTEUR
; APPLICANT: CENTRE NATIONAL DE LA RECHERCHE SCIENTIFIQUE
; TITLE OF INVENTION: VECTORS FOR MOLECULE DELIVERY TO CD11b EXPRESSING CELLS
; FILE REFERENCE: B4669A -AD/DBO
; CURRENT APPLICATION NUMBER: US/10/387,486
; CURRENT FILING DATE: 2003-03-14
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Best Local S
Matches 4
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CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 60/245,755
PRIOR FILING DATE: 2000-11-03
NUMBER OF SEQ ID NOS: 194
SOFTWARE: Patentin version 3.1
SEQ ID NO 38
LENGTH: 11
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: TAN, YI

TITLE OF INVENTION: PRODUCTION OF MOTIF-SPECIFIC AND CONTEXT-INDEPENDENT ANTIBODIES
TITLE OF INVENTION: PEPTIDE LIBRARIES AS ANTIGENS
FILE REFERENCE: CST-138 CIP2
CURRENT APPLICATION NUMBER: US/10/014,485A
CURRENT FILING DATE: 2002-03-18
PRIOR APPLICATION NUMBER: US 09/148,712
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: US 09/535,364
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NUMBER: US 09/535,364
NUMBER OF SEQ ID NUMBER: US 09/535,364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Cell Signaling Technology, Inc. APPLICANT: COMB, Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.1
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NAME/KEY: MOD_RES
LOCATION: (5)..(5)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: (5)...(5)
OTHER INFORMATION: PHOSPHORYLATION; serine at position 5 is phosphorylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
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TYPE: PRT
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80.0%;
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Pred. No. 7.6e+02;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                     1.2e+06;
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; OTHER INFORMATION: Description of Artificial Sequence: intergenic; OTHER INFORMATION: region US-10-387-486-10
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PRIOR FILING DATE: 2000-09-15
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 63, Application US/09879957
Patent No. US20020034755A1
GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
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Best Local (
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PRIOR FILING DATE: 2001-09-14
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                                                                                                                 TELEFAX: (212) 869-81
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 63:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/879,957
FILING DATE: 13-Jun-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION UNMBER: US 08/630,915
APPLICATION NUMBER: US 08/630,915
FILING DATE: 03-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-174
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
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STRANDEDNESS: <Unknown>
TOPOLOGY: unknown
TULE TYPE: peptide
ENCE DESCRIPTION: SEQ ID NO:
                                                                                                  LENGTH: 9 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SPARKS, Andrew B. HOFFMAN, No. US20020034755A1h
                                                                               amino acid
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FOWLKES, Dana M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New York
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71.4%;
                                                                                                                                                                             869-8864/9741
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Pred. No. 1
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US-09-879-957-63

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RESULT 34
US-10-460-512-28
; Sequence 28, Application US/10460512
: Publication No. US20040038271A1
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Best Local Similarity
"httches 4; Conserva
                                       ; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo Sapiens
US-10-460-512-28
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US-10-460-512-6
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 9
 Query Match
                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 28
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Best Local Similarity
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                                                                                                                                                                        APPLICANT: Aya Jakobovits
APPLICANT: Arthur B. Raitano
TITLE OF INVENTION: NOVEL C-TYPE LECTIN TRANSMEMBRANE
TITLE OF INVENTION: ANTIGEN EXPRESSED IN HUMAN PROSTATE CANCER AND USES THEREOF
FILE REFERENCE: 129.20USUI
CURRENT APPLICATION NUMBER: US/10/460,512
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: US/09/638,203
PRIOR APPLICATION NUMBER: 00/148,935
PRIOR APPLICATION NUMBER: 60/148,935
PRIOR APPLICATION NUMBER: 1999-08-12
THEORY OF SECULE 1999-08-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Arthur B. Raitano
TITLE OF INVENTION: NOVEL C-TYPE LECTIN TRANSMEMBRANE
TITLE OF INVENTION: ANTIGEN EXPRESSED IN HUMAN PROSTATE CANCER AND USES THEREOF
FILE REFERENCE: 129.20USU1
CURRENT APPLICATION NUMBER: US/10/460,512
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: US/09/638,203
PRIOR APPLICATION NUMBER: 60/148,935
PRIOR APPLICATION NUMBER: 60/148,935
PRIOR FILING DATE: 1999-08-12
                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                equence 6, Application US/10460512 blication No. US20040038271A1
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 45.1%;
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Pred. No. 1.2e+06;
1; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 23; DB 12; Length 9;
Pred. No. 1.2e+06;
1; Mismatches 2; Indels
 Score 23;
   DB
 12;
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Length 9;
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RESULT 36
US-10-807-856-63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 63, Application to, 1020040157216A1
Publication No. US20040157216A1
CENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11, Application US/10003983B Publication No. US20030103946A1 GENERAL INFORMATION:
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Matches 4; Conservative
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CURRENT FILING DATE: 2001-10-31
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Imperial College Innovations
APPLICANT: Stauss, Hans Josef
APPLICANT: Amrolia, Persis Jal
TITLE OF INVENTION: Immunotherapeutic Methods and Molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
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                                                                                                                       COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, VG
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
hes 4; Conserv
                                                 APPLICATION NUMBER: US/10/807,856
FILING DATE: 23-Mar-2004
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF
ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SPARKS, Andrew B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 YKNSKF 9
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               APPLICATION NUMBER: US/08/630,915 FILING DATE: 03-APR-1996
                                                                                                                                                                                                                                                                          STATE: New York
                                                                                                                                                                                                                                                                                              CITY: New York
                                                                                                                                                                                                                                                                                                              ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/10807856
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INVENTION: POLYPEPTIDES HAVING A FUNCTIONAL
DOMAIN OF INTEREST AND METHODS OF IDENTIFYING AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                            HOFFMAN, Noah
KAY, Brian K.
FOWLKES, Dana M.
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1; Mismatches 2;
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Pred. No. 1.2e+06
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GENERAL INFORMATION:

APPLICANT: Thomas, George
APPLICANT: Kozma, Sara
ITITLE OF INVENTION: P70-S6K or Related Kinase With
ITITLE OF INVENTION: Constitutive Activity
FILS REFERENCE: 4-20793/A/PCT
CURRENT APPLICATION NUMBER: US/10/317,550
CURRENT FILING DATE: 2002-12-12
PRIOR APPLICATION NUMBER: US/09/297,404B
PRIOR FILING DATE: 2002-06-04
NUMBER OF SEQ ID NOS: 24
SEQ ID NO 2
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US-09-572-270A-623
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Publication No. US20030148368A1
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INFORMATION: Inter- complementary peptide listing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/09/572,270A
CURRENT FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 1144
SOFTWARE: ProtPatent version 1.0
SEQ ID NO 623
                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/10317550 Publication No. US20030166034A1
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Best Local :
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TYPE: PRT
ORGANISM: Arabidopsis Thaliana
OTHER INFORMATION: Sequence located in Unknown at 192-201 and may interact with
07-572-270A-623
ORGANISM: Oryctolagus cuniculus
10-317-550-2
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                                        LENGTH: 10
TYPE: PRT
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MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 1101-174 TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Conservative
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Similarity 66.7%;
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(212) 8
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Pred. No. 1.2e+06;
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Friedrich, Thomas

APPLICANT: Zimmerman, Norbert

APPLICANT: Sturmer, Rainer

TITLE OF INVENTION: Reaction of (di)amines in the presence

FITLE OF INVENTION: of a lysin oxidase and of a reducing ager

FILE REFERENCE: BGI-156US

CURRENT APPLICATION NUMBER: US/10/474,601

CURRENT APPLICATION NUMBER: D502/03873

PRIOR APPLICATION NUMBER: PCT/EP02/03873

PRIOR FILING DATE: 2003-10-09

NUMBER OF SEQ ID NOS: 12

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 12
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APPLICANT: Ballinger, Marcus
APPLICANT: Kavanaugh, Michael
TITLE OF INVENTION: Synthetic Peptides Having FGF Receptor
TITLE OF INVENTION: Affinity
FILE REFERENCE: 035784/263218
CURRENT APPLICATION NUMBER: US/10/302,100B
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: 09/407,687
PRIOR APPLICATION NUMBER: 09/407,687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/102,667
PRIOR FILING DATE: 1998-09-30
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 28
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Best Local Similarity
Matches 4; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                             TYPE: PRT ORGANISM: Pichia pastoris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                   LOCATION: 8
OTHER INFORMATION:
                                                                                                              NAME/KEY: VARIANT
                  VAME/KEY: VARIANT
INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 YKNSKF 9
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No. US20040054131A1
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Xaa =
                                                                      Xaa = Any Amino Acid, probably Ser
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66.7%;
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Pred. No. 1.1e+03;
0; Mismatches 2
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Pred. No.
Glu
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1.3e+03;
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LOCATION: (8)...(8)
COTHER INFORMATION: PHOSPHORYLATION; serine at position 8 is phosphorylated
US-10-174-105A-153
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                                                                                                                                                                                                                                                                                                                                       Sequence 919, Application US/10351641 Publication No. US20030186874A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.1 SEQ ID NO 153
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APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: PROPERTIES
FILE REFERENCE: 7872-100
CURRENT APPLICATION NUMBER: US/10/351,641
CURRENT FILING DATE: 2003-01-24
PRIOR APPLICATION NUMBER: 09/350,641
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: 09/315,304
PRIOR APPLICATION NUMBER: 09/315,304
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR FILING DATE: 1998-05-20
                                                                                                                                                                                                                                                                          APPLICANT: Barney, S. APPLICANT: Guthrie, K. APPLICANT: Merutka, G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/535,364
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: CST-138 CIP3
CURRENT APPLICATION NUMBER: US/10/174,105A
CURRENT FILING DATE: 2002-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Cell Signaling Technology, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/148,712 PRIOR FILING DATE: 1998-09-04
                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
OTHER INFORMATION: Synthetic Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPLICANT: COMB, Michael J.

PPLICANT: TAN, Yi

ITLE OF INVENTION: POSITIVE IDENTIFICATION OF PHOSPHO-PROTEINS USING MOTIF-SPECIFIC.

ITLE OF INVENTION: CONTEXT-INDEPENDENT ANTIBODIES COUPLED WITH DATABASE SEARCHING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH: 13
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Pred. No. 1.4e+03;
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US-09-799-250-235
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US-09-799-250-154
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens
US-09-799-250-154
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, Publication No. US200
, GENERAL INFORMATION:
, APPLICANT: Pia M. Ch
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Best Local Similarity 75.0
Thehes 3; Conservative
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                                                                                                                                                                                                                                                                                               Sequence 235, Publication No.
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 919
LENGTH: 8
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Best Local Similarity
SEQ ID NO 235
                                TITLE OF INVENTION: 121P1F1: A TISSUE SPECIFIC PR
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS
FILE REFERENCE: 129.34US01
CURRENT APPLICATION NUMBER: US/09/799,250
CURRENT FILING DATE: 2001-02-08
NUMBER OF SEQ ID NOS: 719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/799,250
CURRENT FILING DATE: 2001-02-08
NUMBER OF SEQ ID NOS: 719
                                                                                                                                            APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mitchell
APPLICANT: Arthur B. Raitano
APPLICANT: Mary Faris
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
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TITLE OF INVENTION: 121P1F1: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCER
FILE REFERENCE: 129.34USO1
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Daniel E.H. Afar
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nilarity 42.9%;
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Pred. No. 1.2e+06;
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APPLICANT: Pia M. Challita-Eid
APPLICANT: Rene S. Hubert
APPLICANT: Steve Chappell Mitchell
APPLICANT: Arthur B. Raitano
APPLICANT: Mary Faris
APPLICANT: Mary Faris
APPLICANT: Mary Faris
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovits
TITLE OF INVENTION: 121P1F1: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: 121P1F1: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN VARIOUS CANCERS
FILE REFERENCE: 129.34US01
CURRENT APPLICATION NUMBER: US/09/799,250
CURRENT FILING DATE: 2001-02-08
NUMBER OF SEQ ID NOS: 719
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 330
LENGTH: 9
TYPE: PRT
CRGANISM: Homo sapiens
US-09-799-250-330
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-799-250-235
Search completed: August 30, 2004, 11:04:55 Job time: 13.0068 secs
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3 FGFEENK 9
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Perfect score:
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| | 7, Appli | 31, Appl | 9 | • | - | 919, App | - | 13, Appl | 30, App1 | ` | • | 50, Appl | | | ` | 5217869 | 28, App1 | |

ALIGNMENTS

; LENGTH: 12 ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-347-926-13 밁 Ś US-09-347-926-13 Sequence 13, Application Sequence 13, Application 13, Application 1440386 Patent No. 6440386 GENERAL INFORMATION: SHUI-ON STABI) CURRENT APPLICATION NUMBER: US/09/347,926 CURRENT FILING DATE: 1999-07-06 NUMBER OF SEQ ID NOS: 24 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 13 LENGTH: 12 Best Loc Matches Query Match Best Local S APPLICANT: LEUNG, SHUI-ON TITLE OF INVENTION: STABILIZED RADIOPHOSPHATE-LABELED FILE REFERENCE: 018733/0936 l Similarity 5; Conserva GFGYKNSKF 9 GESYTNPEF 12 Application US/09347926 Conservative Score 30; Pred. No. Mismatches 4 Length 12 Indels PROTEINS 0; Gaps

0

RESULT 2 US-08-320-373-76

Sequence Patent No

5, Application US/08320373 5559025

Patent No. 5559025 GENERAL INFORMATION:

APPLICANT: APPLICANT: APPLICANT:

Maurer-Fogy, Ingrid Sommergruber, Wolfgang

Ahorn, Horst

APPLICANT: Liebig, Hans-Diet APPLICANT: Skern, Timothy TITLE OF INVENTION: ExpressiTITLE OF INVENTION: Partial TITLE OF INVENTION: Having a NUMBER OF SEQUENCES: 91

Expression of Mature Partial Purification Having an Inhibitory

Proteinase :
 Thereof and
 Effect

and

2A, the d Preparation

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Substrates

CORRESPONDENCE ADDRESS

ADDRESSEE: Sterne, Kessler, Goldstein & STREET: 1225 Connecticut Avenue, N.W. CITY: Washington STATE: D.C. COUNTRY: USA

APPLICANT:
APPLICANT:

Zophel, A...

Blaas, Dieter

Ruchler, Ernst

Liebig, Hans-Dieter

Skern, Timothy

Expression

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Best Local (
                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER9
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 248636 SSK
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
INFORMATION FOR
                                                                                                                                                                                                                                                                                                      ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tent No.
                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/971,619
FILING DATE: 06-NOV-1992
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 466-0800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Oppenheim, Frank G.
APPLICANT: Xu, Tao
APPLICANT: Spacciapoli, Peter
APPLICANT: Roberts, F. D.
APPLICANT: Friden, Philip M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy
                                                                                                                                                                                       APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1996 CLASSIFICATION: 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/320,373
FILING DATE: 11-OCT-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                       E: Hamilton, Brook, Smith & Reynolds, P.C.
Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roberts, F. D.
Friden, Philip M.
VENTION: Anti-Fungal D-Amino Acid Histatin-Based
                                                                                                                                                                                                                                                                                                                                                                      SD
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                                                                                                                                                       US 08/485,273
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                                                                 PER95-02A2
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Pred. No. 3e+05;
2; Mismatches
                                                                                                                                                                                                                                                             Version #1.30
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                                                                                                                   INFORMATION FOR SEQ ID NO: 28
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
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Query Match
Best Local Similarity
Matches 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/973,559

FILING DATE: 07-JUN-1996

CLASSIFICATION: 514

PRIOR APPLICATION NUMBER: US 08/481,888

ENTITY DATE: 07-JUN-1996

CLASSIFICATION NUMBER: US 08/481,888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 28,
                                                                                    TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                            REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-01A2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
TELEPAX: 781-861-9540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                            FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: Region LOCATION: 1..12 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
ZIP: 021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: Two Mil:
CITY: Lexington
                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : YEOLIOGY:
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SPACCIAPOLI, PETER
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                                                                                                   linear
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                                                                                 peptide
                    52.9%;
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    Score 27; DB 2
Pred. No. 78;
0; Mismatches
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                                    Length 12;
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FGYKNSKF 9

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GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. DONALD
APPLICANT: XU, TAO
APPLICANT: XU, TAO
APPLICANT: SPACCIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT APPLICATION NUMBER: US/08/993,235
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 8
LENGTH: 12
TYPE: PRT
ORGANISM: Homo sapiens
US-08-993-235-8
                                                                                                                                                                                                                                                                                                       APPLICANT: Spacciapoli, Peter
APPLICANT: Rothstein, David M.
APPLICANT: Friden, Phillip M.
TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
CURRENT APPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
            RESULT 7
US-08-993-235-8
; Sequence 8, Application US/08993235
; Patent No. 6531573
; GENERAL INFORMATION:
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US-09-226-666-4
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Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 52.9%;
Best Local Similarity 62.5%;
Matches 5; Conservative
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APPLICANT: OPPENHEIM,
                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Derived from Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       No. 6084064
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                                                                                                                                        FGYKRKFF 12
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                                                                                                                                                                                                                      52.9%;
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                                                                                                                                                                                                        Score 27; DB 4;
Pred. No. 78;
0; Mismatches
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APPLICANT: SPACCIAPOLI, PETER
; APPLICANT: SPACCIAPOLI, PETER
; TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
; FILE REFERENCE: 50032/002001
; CURRENT APPLICATION NUMBER: US/08/993,235
; CURRENT FILING DATE: 1997-12-18
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 8
; SEQ ID NO 8
; LENGTH: 12
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-993-235-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Mayfield, Stephen
TITLE OF INVENTION: RNA BINDING PROTEIN AND
FILTE OF INVENTION: EXPRESSION OF RECOMBINA
FILE REFERENCE: SCR21778
CURRENT APPLICATION NUMBER: US/09/613,182
CURRENT FILING DATE: 2000-07-10
PRIOR APPLICATION NUMBER: 09/341,550
PRIOR APPLICATION NUMBER: PCT/US98/00840
PRIOR APPLICATION NUMBER: PCT/US98/00840
PRIOR APPLICATION NUMBER: 60/035,955
PRIOR APPLICATION NUMBER: 60/035,955
PRIOR APPLICATION NUMBER: 60/035,955
PRIOR APPLICATION NUMBER: 60/069,400
PRIOR FILING DATE: 1997-12-12
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Thickes 5; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT; ORGANISM: Chlamydomonas reinhardtii US-09-613-182-2
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US-09-613-182-2
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                                                                                                                                                                                                              RESULT 9
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Best Local Similarity 62.5
                                                                                                                                         Sequence 15, Application US/09649063 Patent No. 6600022 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09613182 Patent No. 6294653
                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                 KUNIKATA, Toshio
KURIMOTO, Masashi
TITLE OF INVENTION: INTERLEUKIN-18 RECEPTOR PROTEINS
NUMBER OF SEQUENCES: 31
                                                                                                     APPLICANT: TORIGOE, USHIO, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ហ
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                                                                                                                                                                                                                                                                                                                                          Conservative
CE ADDRESS:
SE: BROWDY AND NEIMARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        52.9%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROTEIN AND BINDING SITE OF RECOMBINANT MOLECULES
                                                                                                                                                                                                                                                                                                                                          Mismatches
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91;
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419 Seventh Street, N.W., Suite 300

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RESULT 10
US-09-150-621-60
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FILE REFERENCE: 331949-140798
CURRENT APPLICATION NUMBER: US/09/150,621
CURRENT FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: PCT/IL97/00087
PRIOR FILING DATE: 1997-03-10
PRIOR APPLICATION NUMBER: 117426
PRIOR APPLICATION NUMBER: 117426
PRIOR FILING DATE: 1996-03-10
NUMBER OF SEQ ID NOS: 61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                    APPLICANT: YU-CHEN, CHEN
APPLICANT: MUHLRAD, ANDRAS
APPLICANT: SHTEYER, ARIE
APPLICANT: CHOREV, MICHAEL
TITLE OF INVENTION: SYNTHETIC PEPTIDES AND PSEUDOPEPTIDES HAVING OSTEOGENIC
TITLE OF INVENTION: ACTIVITY AND PHARMACEUTICAL COMPOSITIONS CONTAINING THE
TITLE OF INVENTION: SAME
TITLE OF INVENTION: SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/649,063
FILING DATE: 29-Aug-2000
PRIOR APPLICATION NUMBER: 08/996,140
FILING DATE: <Unknown>
APPLICATION NUMBER: JP 52,526/1997
FILING DATE: 21-FEB-1997
APPLICATION NUMBER: JP 163,490/1997
FILING DATE: 6-JUN-1997
APPLICATION NUMBER: JP 215,490/1997
FILING DATE: 28-JUL-1997
APPLICATION NUMBER: JP 215,490/1997
FILING DATE: 28-JUL-1997
APPLICATION NUMBER: JP 215,490/1997
FILING DATE: 28-JUL-1997
APTORNEY/AGENT INFORMATION:
NAME: BROWDY, ROSET L.
REGISTRATION NUMBER: 25,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal fragment
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
RMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 'TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              60, Application US/09150621
5. 6479460
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                                                                          UMBER: PCT/IL97/00087
1997-03-10
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,563A
FILING DATE: 07-JUN-1996
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-JUN-1995
AFTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-02A2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 11
US-08-973-563A-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: peptide US-09-150-621-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
Query Match
Best Local Similarity '5.0
Conservative
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Best Local Similarity 100.

Matches 4; Conservative
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: INC
                                                                                                       NAME/KEY: Region LOCATION: 1..12 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                     OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                   STRANDEDNESS:
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IVENTION: Anti-Fungal D-Amino Acid Histatin-Based
IVENTION: Peptides
REQUENCES: 37
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Roberts,
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                                                                                                                                                                                                       linear
                                                                                                                                                                                     peptide
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                                       48.0%;
75.0%;
                                                                                                     /note= "At least one amino acid must have a D configuration."
                       Score 24.5; DB 2; Length 12; Pred. No. 2.2e+02; 0; Mismatches 1; Indels
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                                                             RESULT 13
US-08-973-559-25
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                              Sequence 25, A Patent No. 591
                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/485,273

FILING DATE: 07-JUN-1995

ATTORNEY/AGENT INFORMATION:

NAME: Brook, David E.

REGISTRATION NUMBER: 22,592

REGISTRATION NUMBER: 2,592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDITED TWO ... STREET: TWO ... CITY: Lexington CTMATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: PETELECOMMUNICATION INFORMATION: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PPLICANT: Roberts, F. D.

IPPLICANT: Roberts, F. D.

IPPLICANT: Roberts, F. D.

IPPLICANT: Roberts, F. D.

IPPLICANT: Philip M.

ITITLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based

TITLE OF INVENTION: Peptides

TITLE OF CHOMENCES: 37
                                                                                                                                                                                                                                                                  NAME/KEY: Region
LOCATION: 1..12
OTHER INFORMATION:
OTHER INFORMATION:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: M
                                                                                                                                                                                       Local Similarity 75.0 tes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 07-JUN CLASSIFICATION: 514
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                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                       NO.
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               INFORMATION
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                                              Application
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Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                      12 amino acids
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OPPENHEIM, FRANK G.
                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                   peptide
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                                                                                                                                                                                                         48.0%;
75.0%;
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must have a D configuration."
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                                              US/08973559
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Pred. No. 2.
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RESULT 14 US-08-973-559-27

Application US/08973559

Sequence 27 Patent No.

GENERAL INFORMATION:

APPLICANT:

OPPENHEIM, FRANK G. XU, TAO

ROBERTS,

F.D.

APPLICANT: SPACCIAPOLI, PETER
APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:

STREET: TWO ... CITY: Lexington OTATE: MA

ADDRESSEE: Hamilton, Brook, Smith & Reynolds, STREET: Two Militia Drive

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

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Query Match
Best Local Similarity
"Arches 6; Conserva
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                                                                                                                                US-08-973-559-25
                                                                                                                                                                                                                                                                                                                           FILING DATE: 07-JUN-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0,

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/973,559
                                                                                                                                                                                                          TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SPACCIAPOLI, PETE APPLICANT: FRIDEN, PHILIP M. TITLE OF INVENTION: Anti-Fun TITLE OF INVENTION: Histatin NUMBER OF SEQUENCES: 37
                                                                                                                                              MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: PE
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02173
                                                                                                                                                                                             amino acid
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 FGYKR-KF 11
                               FGYKNSKF 9
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                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ROBERTS, F. D.
                                                                                                                                                               linear
                                                                                                                                                                                                                                                                               781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti-Fungal and Anti-Bacterial
Histatin-Based Peptides
: 37
                                                                              48.0%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Release #1.0, Version #1.30
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                                                                              Score 24.5;
Pred. No. 2.
                                                                Mismatches
                                                                               .2e+02
                                                                                                 DB
                                                                                             Length 12;
                                                                  Indels
                                                                1;
                                                                Gaps
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APPLICANT: OPPENHEIM, FRANK G.

APPLICANT: ROBERTS, F. DONALD

APPLICANT: XU, TAO

APPLICANT: SPACCIAPOLI, PETER

TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES

FILE REFERENCE: 50032/002001

CURRENT APPLICATION NUMBER: US/08/993,235

CURRENT FILING DATE: 1997-12-18

NUMBER OF SEQ ID NOS: 13

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 5

LENGTH: 12

TYPE: PRT

ORGANISM: Homo sapiens

US-08-993-235-5
                                                                                                                                                                                                                          Query Match .
Best Local Similarity
Marches 6; Conserva
                                                                                                     RESULT 16
US-08-993-235-7
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US-08-993-235-5
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Sequence 7, Application US/08993235
Patent No. 6084064
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. DONALD
APPLICANT: XU, TAO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION.

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/481,888

FILING DATE: 07-JUN-1995

ATTORNEY/AGENT INFORMATION:

NAME: Brook, David E.

NAME: Brook, David E.

PER95-01A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5, Application US/08993235 Patent No. 6084064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide -08-973-559-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PE
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08
FILING DATE: 07-JUN-1996
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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Similarity 75.0%;
6; Conservative
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Pred. No. 2.2e+02;
0; Mismatches 1;
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; APPLICANT: SPACCIAPOLI, PETER
; TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
; FILE REFERENCE: 50032/002001
; CURRENT APPLICATION NUMBER: US/08/993,235
; CURRENT FILING DATE: 1997-12-18
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 7
; LENGTH: 12
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-993-235-7
                                                                                                                                                                                                                                                                                                                          RESULT 18
US-08-993-235-7
; Sequence 7, Appl
; Patent No. 65319
; LENGTH: 12
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-993-235-7
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US-08-993-235-5
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                                                                           NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
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TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT APPLICATION NUMBER: US/08/993,235
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 5
FENCING 10
                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. DONALD
APPLICANT: XU, TAO
APPLICANT: SPACCIAPOLI, PETER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 6; Conserv
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Best Local Similarity 75.0%;
Matches 6; Conservative
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Patent No. 653157
                                                                                                                                        CURRENT APPLICATION NUMBER: US/08/993,235 CURRENT FILING DATE: 1997-12-18
                                                                                                                                                                              APPLICANT: SPACCIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL
FILE REFERENCE: 50032/002001
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TYPE: PRT
ORGANISM: Homo sapiens
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Pred. No. 2.2e+02;
0; Mismatches 1
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Pred. No. 2.2e+02;
0; Mismatches 1;
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RESULT 20
5217869-104
;Patent No. 5217869
; Patent No. 5217869
; APPLICANT: KAUVAR, LAWRENCE M.
TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                          RESULT 21
US-08-135-319A-8
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APPLICANT: Roggen, Erwin
APPLICANT: Roggen, Erwin
APPLICANT: Ernst, Steffen
TITLE OF INVENTION: Low Allergenic Protein Variants
FILE REFERENCE: 5676.200-US
CURRENT APPLICATION NUMBER: US/09/417,608A
CURRENT FILING DATE: 1999-10-13
NUMBER OF SEQ ID NOS: 101
SOFTWARE: PatentIn version 3.2
SEQ ID NO 86
LENGTH. 0
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US-09-417-608A-86
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Best Local S
Matches 4
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                                     GENERAL INFORMATION:
APPLICANT: Heavner, George A.
APPLICANT: McEver, Rodger P.
APPLICANT: Geng, Jian-Guo
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Best Local
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Best Local (
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APPLICANT: Geng, Jian-Guo
TITLE OF INVENTION: Peptide Inhibitors of Inflammation Mediated by Selectins
FILE REFERENCE: CTC 102 CON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 121
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
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Pred. No. 3e+0
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Pred. No. 3e+05;
1; Mismatches
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Pred. No. 2.2e+02;
0; Mismatches 1
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Query Match
Best Local Similarity
Thickes 4; Conserva
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US-09-347-926-9
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Sequence 9, Application US/09347926
Patent No. 6440386
GENERAL INFORMATION:
APPLICANT: LEUNG, SHUI-ON
TITLE OF INVENTION: STABILIZED RADIOPHOSPHATE-LABELED PROTEINS
FILE REFERENCE: 018733/0936
CURRENT APPLICATION NUMBER: US/09/347,926
CURRENT FILING DATE: 1999-07-06
NUMBER OF: SEQ ID NOS: 24
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 9
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CURRENT FILING DATE: 1993-10-12
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
SEQ ID NO 8
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Patent No. 6624295
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Best Local S
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CURRENT FILING DATE: 1999-08-26
EARLIER APPLICATION NUMBER: US 60/098,233
EARLIER FILING DATE: 1998-08-28
EARLIER APPLICATION NUMBER: US 60/122,767
EARLIER FILING DATE: 1999-03-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Adams, Camelia W.
APPLICANT: Devaux, Brigitte
APPLICANT: Eaton, Dan L.
APPLICANT: Hass, Philip E.
APPLICANT: Judice, J. Kevin
APPLICANT: Kirchhofer, Daniel
APPLICANT: Suggett, Shelley
TITLE OF INVENTION: Human Anti-Factor IX/IXa Antibodies
FILE REFERENCE: P1661R2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 32
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ORGANISM: artificial sequence
FEATURE:
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4; Conservative
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Pred. No.
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Pred. No. 2.5e+02;
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ORGANISM: Homo sapiens

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RESULT 25
US-08-215-805A-3
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5217869-103
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APPLICANT: KAUVAR, LAWRENCE M.

TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
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Best Local S
Matches 4
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LENGTH: 9
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Best Local .
                                                                                         TELEFAX: (716) 263-1600 INFORMATION FOR SEQ ID NO: 3:
                                                                                                                   NAME: Timian, Susan J.
REGISTRATION NUMBER: 34,103
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
TELEPHONE: (716) 263-1636
                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 121
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                            APPLICATION NUMBER: US/08
FILING DATE: 22-MAR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                              CITY: Rochester
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                               STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION:
DANT: Chang, Yung-Fu
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                                            amino acid
TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                         E: Nixon, Hargrave, Devans & Doyle Clinton Square, P.O. Box 1051
                                                                                                                                                                                                                                                                                                                                                                USA
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                linear
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85.7%;
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                                                                                                                                                   19603/61 (D-1329A)
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Pred. No. 3e
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Pred. No. 2.
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                                                                                                                                                                                                                                                                        Version #1.30
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3e+05;
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RESULT 27
US-08-630-915A-63
; Sequence 63, Application US/08630915A
; Patent No. 6309820
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US-08-215-805A-41
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                                                                                                                                                              Query Match
Best Local Similarity
Matches 4; Conserv
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Patent No. 5559008
GENERAL INFORMATION:
APPLICANT: Chang, Yung-Fu
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INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                            MOLECULE TYPE: peptide ORIGINAL SOURCE: ORGANISM: Actinobaci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/215,805A
FILING DATE: 22-MAR-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Timian, Susan J.
REGISTRATION NUMBER: 34,1
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Chang, Yung-Fu
TITLE OF INVENTION: LEUKOTOXIN GENE FROM PASTEURELLA
TITLE OF INVENTION: SUIS
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IMMEDIATE SOURCE:
LIBRARY: P. suis DNA in Bacteriophage lambda-dash
CLONE: (Lambda)yfc33-37
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ORGANISM: Pas
                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                       TOPOLOGY:
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T: Clinton Square, P.O. Box 1051
Rochester
: New York
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66.7%;
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66.7%;
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Pred. No. 3e+05;
1; Mismatches
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Pred. No. 3e+05;
1; Mismatches
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US-09-638-203-6
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ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,915A
FILING DATE: 03-APR-1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 1101-174
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
TELES: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 63:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: Amino acids
SOFTWARE:
SEQ ID NO 6
LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                      Sequence 6, Application US/09638203 Patent No. 6602501 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -08-630-915A-63
                                                            APPLICANT: Daniel E.H. Afar APPLICANT: Rene S. Hubert APPLICANT: Aya Jakobovits APPLICANT: Arthur B. Raitanc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SPARKS
APPLICANT: HOFFMAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: KAY, Brian K.
APPLICANT: FOWLKES, Dana M.
APPLICANT: McCONNELL, Stephen J.
TITLE OF INVENTION: POLYPEPTIDES HAVING A FUNCTIONAL
TITLE OF INVENTION: DOMAIN OF INTEREST AND METHODS OF IDENTIFYING
TITLE OF INVENTION: USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: ur
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STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       45.1%;
Similarity 66.7%;
4; Conservation
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                                                  FastSEQ for Windows Version 4.0
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HOFFMAN, No. 63098
KAY, Brian K.
FOWLKES, Dana M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unknown
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Pred. No. 3e+05;
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APPLICANT: Daniel E.H. Afar

APPLICANT: Rene S. Hubert

APPLICANT: Aya Jakobovits

APPLICANT: Aya Jakobovits

APPLICANT: Arthur B. Raitano

TITLE OF INVENTION: NOVEL C-TYPE LECTIN TRANSMEMBRANE

TITLE OF INVENTION: ANTIGEN EXPRESSED IN HUMAN PROSTATE CANCER AND USES THEREOF

FILE REFERENCE: 129.20USU1

CURRENT APPLICATION NUMBER: US/09/638,203

CURRENT FILING DATE: 2000-08-11

PRIOR APPLICATION NUMBER: 60/148,935

PRIOR APPLICATION NUMBER: 60/148,935

PRIOR FILING DATE: 1999-08-12

NUMBER OF SEQ ID NOS: 47

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 28 \

TYPE: PRT

ORGANISM: Homo Sapiens

US-09-638-203-28
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US-09-638-203-28
; Sequence 28, Ap
                                                                                                                                                Query Match
Best Local Similarity
Thickes 4; Conserve
                                            RESULT 31
US-08-855-958-4
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Best Local Similarity
"atches 4; Conserva
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                                                                                                                                                                                                                                                                                                                                     ;Patent No. 5217869
; APPLICANT: KAUVAR, L
; TITLE OF INVENTION:
;REAGENTS
                                                                                                                                                                                                                                                          SEQ ID NO:115:
Sequence 4, Application US/08855958B Patent No. 6143723 GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 121 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                              LENGTH: 9
                                                                                                 1 GFGKKHT 7
                                                                                                                                    1 GFGYKNS 7
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57.1%;
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57.1%;
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Pred. No. 3e+0
1; Mismatches
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Pred. No. 3e+05;
2; Mismatches
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Pred. No. 3e+05;
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3e+05;
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; Sequence 28, Application US/09407687
; Patent No. 6548634
; GENERAL INFORMATION:
APPLICANT: Ballinger, Marcus
APPLICANT: Kavanaugh, Michael
TITLE OF INVENTION: Synthetic Peptides Having FGF Receptor
TITLE OF INVENTION: Affinity
FILE REFERENCE: 1517.001
; CURRENT APPLICATION NUMBER: US/09/407,687
; CURRENT FILING DATE: 1999-09-28
; EARLIER APPLICATION NUMBER: 60/102,667
; EARLIER FILING DATE: 1998-09-30
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US-08-855-958-9
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; OTHER INFORMATION: cyclo peptide
US-08-855-958-9
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SEQ ID NO 9
LENGTH: 11
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Best Local 9
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ramaiah, Abburi
TITLE OF INVENTION: PIGMENTARY AGENT
FILE REFERENCE: 31446-132213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Ramaiah, Abburi
TITLE OF INVENTION: PIGMENTARY AGENT
FILE REFERENCE: 31446-132213
CURRENT APPLICATION NUMBER: US/08/855,958B
CURRENT FILING DATE: 1997-05-14
NUMBER OF SEQ ID NOS: 9
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TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE: OTHER INFORMATION: Description of Unknown Organism: bFGF fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: cyclo peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: Description of Unknown Organism: bFGF fragment
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3; Conserv
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FastSEQ for Windows Version 3.0
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Pred. No.
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Pred. No. 3.9e+02;
3; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 3;
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APPLICANT: AOYAGI, KAORU
APPLICANT: SUMIDA, NAOMI
APPLICANT: MURAKAMI, TAKESHI
APPLICANT: MURAKAMI, TAKESHI
APPLICANT: MURAKAMI, TAKESHI
APPLICANT: MURAKAMI, TAKESHI
APPLICANT: NUVENTION: REGULATORY SEQUENCE OF CELLULASE cbh
TITLE OF INVENTION: IN TRICHODERMA VIRIDE AND SYSTEM FOR
TITLE OF INVENTION: PROTEINS OR PEPTIDES THEREWITH
FILE REFERENCE: 99-0266*/LC(WMC)/00144
CURRENT APPLICATION NUMBER: US/09/254,733
CURRENT FILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 52
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; FEATURE:
; OTHER INFORMATION: synthetic peptide
US-09-407-687-28
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US-09-254-733-50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: TRICHODERMA VIRIDE MC300-1 US-09-254-733-50
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Best Local Similarity
Matches 4; Conserv
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LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                        Sequence 14, Application US/08310429A Patent No. 5695935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 50
LENGTH: 14
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                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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APPLICANT: MORIYA, TATSUKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                              APPLICANT: Page, David C.
APPLICANT: Reijo, Renee
TITLE OF INVENTION: DAZ: A GENE ASSOCIATED WITH AZOOSPERMIA
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
                                                                                           ZIP: 02173
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.30 URRENT APPLICATION DATA: APPLICATION NUMBER: US/08/310,429A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity les 4; Conserv
                                                                                                                                                                   CITY: Lexington
STATE: Massachusetts
                                                                                                                                                                                                        STREET:
                                                                                                                                                  COUNTRY:
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Pred. No. 4.9e+02
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Pred. No. 4.2e+02;
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FILING DATE:

22-SEP-1994

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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                     TELEFAX: (617) 861-954
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Page,
APPLICANT: Reijo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 equence 13, Application US/08690734A atent No. 5871920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                             FILING DATE: 31-JUL-1
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/690,734A
                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: WH
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 861-6240
                                                              TOPOLOGY: li
                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
ATTORNEY/AGENT IN
                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STATE: Massachus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                   NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/310,429 FILING DATE: 22-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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                                                                                                            : 8 amino acids amino acid
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Two Militia Drive
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                                                            peptide
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  43.1%;
50.0%;
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 Score 22;
Pred. No.
                                                                                                                                                                                                                       WHI94-07A
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Pred. No.
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               Length 8;
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SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
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                                                     Matches
                                                                Query Match
Best Local
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: H
TITLE OF INVENTION: T
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
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STREET: La Jolla
STATE: CA
STATE: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                Local Similarity
les 4; Conserv
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ADDRESSEE:
                                                                                                                   LOCATION: 8
OTHER INFORMATION:
                                                                                                                                                NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: UPFILING DATE: 19930315
                                                                                                                                                                        LOCATION: 5
OTHER INFORMATION: /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 92037
                                                                                                                                                                                                                               LOCATION: 2
OTHER INFORMATION: /note=
                                                                                                                                                                                                                                                      NAME/KEY: Modified-site
                                                                                                                                                                                             NAME/KEY: Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                    ELEPHONE:
                                                                                                                                                                                                                                                                                       ECULE TYPE: peptide
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                          GYKNS 7
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10666 No. 5968817th Torrey Pines Road, TPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sutcliffe, J Gregor
Erlander, Mark G
Lovenberg, Timothy W
VENTION: HUMAN SEROTONIN RECEPTORS, DNA ENCODING
VENTION: THE RECEPTORS, AND USES THEREOF
EQUENCES: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                          tting, Thomas
                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                      619-554-2937
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                                                                43.1%;
                                                                                                                      /note=
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                                                  Score 22; DB
Pred. No. 3e+0
0; Mismatches
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RESULT 38 US-08-742-185-13

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Best Local S
Matches 3
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                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Barney,
APPLICANT: Guthrie
                                                                                                                                                                                                                     Sequence 919, Application US/09082279B
                                              APPLICANT: Barney, Shawn
APPLICANT: Guthrie, Kelly
APPLICANT: Merutka, Gene
APPLICANT: Anwer, Mohmed
APPLICANT: Lambert, Dennis
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: PHARMACOKINETIC PROPERTIES
FILE REFERENCE: 7872-043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/690,734
FILING DATE: 31-JUL-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/310,429
FILING DATE: 22-SEP-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                           -09-082-2798-919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617) 861-9540 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
CURRENT APPLICATION NUMBER: US/09/082,279B
CURRENT FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: WHI94-07A2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 861-624
TELEFAX: (617)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook,
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Hawkins,Trevor
APPLICANT: Reeve, Mary Pat
ITTLE OF INVENTION: DAZ: A GENE FAMILY ASSOCIATED WITH AZOOSPERMIA
NUMBER OF SEQUENCES: 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/742,185 FILING DATE: 30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Lexington
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Saxena, Richa
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linear
F:
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                            43.1%;
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Pred. No. 3e+05;
2; Mismatches
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US-09-834-784-919
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US-09-315-304B-919
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GENERAL INFORMATION:
APPLICANT: Barney, S.
APPLICANT: Guthrie, K.
APPLICANT: Merutka, G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Core polypeptide US-09-315-304B-919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: PROPERTIES
FILE REFERENCE: 7872-052
CURRENT APPLICATION NUMBER: US/09/315,304B
CURRENT FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR APPLICATION NUMBER: 09/082,279
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 919
LENGTH: 8
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Best Local Similarity 75.0
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Best Local S
                                                                                                      APPLICANT: Barney, Shawn
APPLICANT: Barney, Shawn
APPLICANT: Guthrie, Kelly
APPLICANT: Merutka, Gene
APPLICANT: Merutka, Gene
APPLICANT: Anwer, Mohmed
APPLICANT: Lambert, Dennis
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: PHARMACOKINETIC PROPERTIES
FILE REFERENCE: 7872-043
FILE REFERENCE: 7872-043
                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                CURRENT APPLICATION NUMBER: US/09/834,784
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR FILING DATE: 1998-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
NUMBER OF SEQ ID NOS: 1515
SOFTWARE: FastSEQ for Windows Version 3.0
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75.0%;
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75.08;
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Pred. No. 3e+05
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Pred. No. 3e+05;
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RESULT 43
US-09-350-641C-919
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; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Core polypeptide
US-09-834-784-919
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Best Local S
Matches 3
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Best Local Similarity 75.0
Thes 3; Conservative
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APPLICANT: Delmedico, M.
APPLICANT: Erickson, J.
APPLICANT: Lambert, D.
APPLICANT: Sista, P.
                                                                                                                                                   APPLICANT: Barney, S. APPLICANT: Guthrie, K APPLICANT: Merutka, G APPLICANT: Anwer, M. APPLICANT: Lambert, D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 919, Application US/09515965A
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LENGTH: 8
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Patent No. 6656906
SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lambert, D.
APPLICANT: Sista, P.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION OF MEMBRANE FUSION-
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
FILE REFERENCE: 7872-073
CURRENT APPLICATION NUMBER: US/09/515,965A
CURRENT FILING DATE: 2000-02-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6623741 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 09/315,304
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1994
SOFTWARE: FastSEQ for Windows Version 3.0
                                   CURRENT APPLICATION NUMBER: US/09/350,641C CURRENT FILING DATE: 1999-07-09 PRIOR APPLICATION NUMBER: 09/315,304
                                                                                            APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: PROPERTIES
FILE REFERENCE: 7872-067
PRIOR APPLICATION NUMBER: 09/315,304 PRIOR FILING DATE: 1999-05-20 PRIOR APPLICATION NUMBER: 09/082,279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Artificial Sequence
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3; Conservative
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Merutka, G.
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75.0%;
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1; Mismatches
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Pred. No. 3e+05;
l; Mismatches
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3e+05;
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SOFTWARE: Pat SOFTWARE: Pat ; SEQ ID NO 7

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APPLICANT: Thorsett, Eugene D
APPLICANT: Yednock, Theodore A
APPLICANT: Pleiss, Michael A
TITLE OF INVENTION: Inhibitors of Leukocyte Adi
FILE REFERENCE: 123-US-CIP1
CURRENT APPLICATION NUMBER: US/08/467,580B
CURRENT FILING DATE: 1995-06-06
EARLIER APPLICATION NUMBER: 08/273,055
EARLIER FILING DATE: 1994-07-11
NUMBER OF SEQ ID NOS: 163
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 31
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US-08-990-015-7
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Sequence 7, Application US/08990015A
Patent No. 6093701
GENERAL INFORMATION:
APPLICANT: Budker, Vladimir
APPLICANT: Budker, Vladimir
APPLICANT: Sebestyen, Magdolna G.
APPLICANT: Sebestyen, Magdolna G.
APPLICANT: Wolff, Jon A.
APPLICANT: Wolff, Jon A.
TITLE OF INVENTION: A METHOD FOR COVALENT ATTACHMENT OF COMPOUNDS TO GENES
CURRENT APPLICATION NUMBER: US/08/990,015A
CURRENT APPLICATION NUMBER: US/08/990,015A
CURRENT FILING DATE: 1997-12-12
EARLIER APPLICATION NUMBER: 60/050842
EARLIER FILING DATE: 1997-05-26
NUMBER OF SEQ ID NOS: 11
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Best Local Similarity
Matches 3; Conserv
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NUMBER OF SEQ ID NOS: 1757
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Peptide OTHER INFORMATION: derived from first domain of VCAM-1
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; TYPE: PRT; ORGANISM: Artificial Sequence; ORGANISM: Artificial Sequence; FEATURE: ; FEATURE: ; OTHER INFORMATION: Description of Artificial Sequence: CONTAINS; OTHER INFORMATION: SEQUENCES FROM THE SV40 GENOME US-08-990-015-7 Ş Query Match 43.1%; Score 22; DB 3; Length 10; Best Local Similarity 66.7%; Pred. No. 5.4e+02; Matches 4; Conservative 0; Mismatches 2; Indels 3 GYKNSK 8 ||| | 2 GYKKRK 7 0; Gaps

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Search completed: August 30, 2004, 10:57:12 Job time : 4.55743 secs

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Result
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Maximum Match 100%
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1: geneseqp1980s:*
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4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*
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Match
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Gapop 10.0 , Gapext 0.5
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51
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   GFGYKNSKF 9
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                                                                                                                                                                                                                                       Length DB
   GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                   AAY69923
AAY69924
ABG70966
ADD44040
ADD44091
AAR72924
AAR72880
AAW33943
ABB82364
AAW19418
                           AAW19464
AAY25144
AAB11978
AAB33942
AAW16616
AAW262420
AAM97164
AAM99539
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Aay69924 Human cyc
Abg70966 Human pro
Add44040 CPG2 pept
Add44091 CPG2 pept
Aar72924 Yeast PPI
Aar72880 Yeast PPI
Aaw33943 Betal-adr
Abb82364 BAK prote
Aaw19418 Anti-fung
Aaw19464 Anti-fung
Aaw19464 Anti-fung
Aaw25144 Antibacte
Aab11978 Histatin-
Aaw68100 Chlamydom
Aaw68100 Chlamydom
                                    Adb71196
Aam99439
  Aaw63824
Aab46262
                 Aaw28520
                            Aam99577
                                                                       Aao22420
                                                                                 Aaw1661
                                                                                                                                                                                                                                      Description
) Human Her
! Human Mab
! HPV type
                           Phosphoin
Protease
Human pep
Vaccine r
Human mod
Vaccine r
Vaccine r
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                          Query Match
Best Local S
Matches 9
                          Similarity
9; Conserve
GFGYKNSKF
                         100.0%; Score 51; DB 3; larity 100.0%; Pred. No. 1.4e+06; Conservative 0; Mismatches 0;
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| | 44 | | 42 | | | 39 | 38 | | | S | 34 | | 32 | 31 | 30 | 29 | ω | 27 | 26 |
|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|-----------|----------|-----------|-----------|-----------|-----------|-----------|-----------|----------|----------|-----------|
| 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 | 25 |
| | 9 | 9 | | 9 | 9. | 49.0 | 9. | | 9. | 9. | 9 | 9 | 9 | 9 | | 9 | 9 | 9 | 9 |
| 14 | 14 | 14 | 14 | 14 | 14 | 14 | 14 | 14 | 14 | 14 | 13 | 13 | 13 | 13 | 12 | 12 | 12 | 12 | 10 |
| J. | σ | σ | v | տ | v | 4 | 4, | 4 | 4, | N | 7 | 7 | ω | N | ហ | ഗ | 4, | 4, | Q |
| | ABR59105 | ABR91220 | ABG67720 | σ | AAE25488 | φ | AAU15641 | AAU25297 | | ADC73226 | | ADD44065 | | AAW94978 | AAE28582 | ABB74314 | AAB81463 | AAB81464 | ABR91208 |
| Ada23853 | ŧπ | 912 | Abg67720 | 5 | Aae25488 | Abb52298 | Aau15641 | Aau25297 | Aau68422 | w | Add44039 | Add44065 | Aab13279 | Aaw94978 | Aae28582 | Abb74314 | Aab81463 | Aab81464 | Abr91208 |
| Alzheimer | Alzheimer | P. papata | Human ADP | Human 125 | CAPI-19 t | Human API | Schizophr | Schizophr | Human Bre | | CPG2 pept | CPG2 pept | Caenorhab | MHC bindi | Desmodium | Simple nu | cheti | æ | P. papata |

ALIGNMENTS

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RESULT 1
AAY69923
ID AAY69923;
XX
AC AAY69923;
XX
AC AAY69923;
XX
AC AAY69923;
XX
DT 11-APR-2000 (first entry)
XX
DT 11-APR-2000 (first entry)
XX
Cyclophilin B; human; tumour antiger
XX
Cyclophilin B; human; tumour; then
XX
Cyclophilin B; human; tumour antiger
XX
Cyclophilin B; human; tumour antigen
PD cyclophilin Cyclophiles
Cyclophiles are tumour antigen peptide
Cyclophiles are used for the treatment
XX
Cyclophilin A; Page 50; 64pp; Japanese.
XX
Cyclophiles are used for the treatment
                                                                                        This sequence represents a cyclophilin B peptide of the invention. The peptides are tumour antigen peptides derived from cyclophilin B, that recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                             Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cyclophilin B; human; tumour antigen peptide; cytotoxic T-cell; CTL; HLA antigen; diagnosis; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human cyclophilin B peptide fragment
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Length 9; Indels

°.

Gaps

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RESULT 2
AAX69924
ID AAY6
XX AAX69924
AC AAY6
AC AAY6
AC AAY6
XX Hum
Cyc.
KW Cyc.
KW HLA
XX HOM
XX HOM
XX HOM
YX WO9
PD 29-
XX 29-
XX (SU
PR 25-
XX (SU
PR UIC
XX WP:
XX WP:
XX WP:
XX C1
XX C2-
XX C2
RESULT 3
ABG70966
ID ABG7
XX
AC ABG7
XX
DT 04-D
XX
DE Huma
XX
Huma
KW Huma
KW Phos
KW path
KW path
KW pyot
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Best Local S
Matches 7
                                                          Human; radiotherapeutic agent precursor; targeting peptide; tumour; phosphorylation peptide; linker; Src homology region 2 domain; SH_2; infectious lesion; breast; colon; rectum; prostate; ovary; testes; pathological lesion; chest; abdomen; pelvis; skin; cardiovascular lesion; myocardial infarct; atherosclerotic plaque; clot; thrombosis; pulmonary embolism; inflammatory lesion; hyperplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents a cyclophilin B peptide of the invention. peptides are tumour antigen peptides derived from cyclophilin B, the recognise cytotoxic T-cells (CTL) which bind to the HLA antigen. The peptides are used for the treatment and diagnosis of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumor antigen peptides derived from cyclophilin B for treatment and diagnosis of tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclophilin B; human; to HLA antigen; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY69924 standard; peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-116932/10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-JUN-1999;
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                                                                                                                                                                                                                                                                                                               Human protein kinase Cbetal peptide.
                                                                                                                                                                                                                                                                                                                                                                               04-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABG70966 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; Page 50; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SUMU )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUMITOMO PHARM CO LTD ITOH K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gomi S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GYKNSKF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GYKNSKF 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           76.5%; Score 39; DB 3; ilarity 100.0%; Pred. No. 2.3; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-JP003360
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s; tumour; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               #4
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RESULT 4
ADD44040
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Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             is useful for preparing radiotherapeutic agents, useful for the treatment of tumour and an infectious lesion. The lesions include infectious lesions of breast, colon, rectum, prostate, ovary and testes; pathological lesions affecting chest, abdomen and pelvis and skin; and cardiovascular lesions such as myocardial infarct, atherosclerotic plaque, clot, thrombosis, pulmonary embolism, inflammatory lesions and hyperplasia. The linker in the radiotherapeutic agent is long and sufficiently flexible to allow the SH 2 domain to make a turn and fold back on the phosphorylated substrate. Thus, the radiotherapeutic agent exhibits improved stability in vivo, by avoiding the ready phosphorylation, while maintaining the binding ability of the agent. The present amino acid sequence represents a human peptide that contains an SH 2 domain as described in the invention
                                                                                                                                                           bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand; immunostimulant; enzyme therapy; immune response; gene directed enzyme prodrug strategy; vaccine; enzyme; EC 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising a protein containing a targeting peptide, a phosphorylation peptide, a linker and Src homology region 2 domain (SH 2). The inventic is useful for preparing radiotherapeutic agents, useful for the treatment of tumour and an infectious lesion. The lesions include infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Radioactive agent precursor, useful in manufacture of in vivo radiotherapeutic agents, comprises protein containing targeting phosphorylation peptide, linker and Src homology region 2 domain
                                                                                                                                                                                                                                           CPG2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12 AA;
29-NOV-2001; 2001EP-00128519
                               27-NOV-2002;
                                                                                                                           Pseudomonas sp.
                                                                                                                                                                                                                                                                                                                                          ADD44040 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a new radiotherapeutic agent precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Col 7; 13pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-722167/78.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Leung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6440386-B1.
                                                               05-JUN-2003
                                                                                               WO2003045426-A1
                                                                                                                                                                                                                                                                           15-JAN-2004
                                                                                                                                                                                                                                         peptide #60 with potential human MHC class II binding activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GFGYKNSKF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                        GESYTNPEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                               2002WO-EP013351
                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                              RS-16.
                                                                                                                                                                                                                                                                                                                                        peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                        12
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                                                                                                                                                                                                                                                                                                                                           13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 30; DB
Pred. No. 1e+0
1; Mismatches
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1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 12;
                                                                                                                                                              enzyme; EC 3.4.17.11.
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CC The invention relates to a novel modified bacterial enzyme CC carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than CC calterations to compared with the same biological specificity when used in vivo, and comprising specific amino acid residues having CC alterations cause a reduction or an elimination of one or more of T-cell cC epitope sequences, which act in the parental enzyme as MHC class II CC binding ligands and stimulate T-cells. The modified CPG2 enzyme and the CPG2 proteins have immunostimulant activity and may be used in enzyme CC therapy. The modified CPG2 enzyme may be used to induce an immune CC response in a human host, or as a therapeutic entity such as the gene CC directed enzyme prodrug strategy. The peptide is useful for the CC manufacture of a modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo. This sequence represents a CPG2 enzyme peptide with potential human MHC CC class II binding activity of the invention.
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Best Local S
Matches 5
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                                                                                                                                                                                                         bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand; immunostimulant; enzyme therapy; immune response; gene directed enzyme prodrug strategy; vaccine; enzyme; EC 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JAN-2002;
13-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; Page 13; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     substantially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hellendoorn K,
                                                                                                                                                                             Pseudomonas sp.
                                                                                                                                                                                                                                                                                                                15-JAN-2004
                                                                                                                                                                                                                                                                                                                                              ADD44091;
                                                                                                                                                                                                                                                                                                                                                                           ADD44091 standard;
                            29-NOV-2001; 2001EP-00128519.
25-JAN-2002; 2002EP-00001778.
13-SEP-2002; 2002EP-00020634.
                                                                                        27-NOV-2002; 2002WO-EP013351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified bacterial enzyme carboxypeptidase G2 stantially non-immunogenic or less immunogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2003-513617/48.
                                                                                                                                                                                                                                                                                   peptide #111 with potential human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful for inducing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
5; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13
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2002EP-00020634
                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                RS-16.
                                                                                                                                                                                                                                                                                                                                                                           peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            58.8%;
62.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     an immune response in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Williams
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Score 30; DB;
; Pred. No. 1.1e.
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 13;
                                                                                                                                                                                                            enzyme; EC 3.4.17.11
                                                                                                                                                                                                                                                                                    II binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                     activity.
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(MERE) MERCK PATENT GMBH

Hayano T,

Katou

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Maki N,

Takahashi N,

Suzuki

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RESULT 6
AAR72924
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel modified bacterial enzyme carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any non-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having calterations compared with the non-modified parochial enzyme. The class of epitope sequences, which act in the parental enzyme as MHC class of compared inguity and stimulate T-cells. The modified CPG2 enzyme and the charapy. The modified CPG2 enzyme may be used to induce an immune cresponse in a human host, or as a therapeutic entity such as the gene directed enzyme prodrug strategy. The peptide is useful for the manufacture of a modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo. This sequence represents a CPG2 enzyme peptide with potential human MHC class II binding activity of the invention.
PRACTOR REPRESENTATION OF A CONTRACT OF A CO
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                                                                                                                            19-JUL-1989;
06-OCT-1989;
29-DEC-1989;
19-JUL-1990;
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25-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Escherichia coli; protein conformation;
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5; Conserv
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                                                                         CORP.
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89JP-00260244.
89JP-00344705.
90EP-00307914.
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Pred. No. 1.1e+02;
1; Mismatches 2
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25-MAR-2003
29-NOV-1995
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AAR72877-85 are tryptic fragments of a yeast PPIase trans isomerase). The yeast PPIase has a single moland a single isoelectric point of about 6.2. The enz
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the folding of proteins, partic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      coli; protein conformation;
a; peptidyl prolyl cis trans
on; prolyl peptide bond.
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1.4e+06;
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isomerase alpha; catalysis;
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This sequence is a betal-adrenergic receptor antagonist, and is an antagonist of the invention. The antagonists (A) are for inhibiting the function of a prokaryotic or eukaryotic integral membrane protein (IMP) having at least one transmembrane domain (TMD), comprises a peptide including at least 4 consecutive amino acids (aa) from the sequence of the TMD. (A) are used to treat or prevent disorders in mammals that involve disturbances of IMP, and the same effect is achieved by inserting a nucleic acid encoding (A) into the cells of the mammal. Specified conditions that can be treated are schizophrenia, psychotic disorders, Huntington's disease, Tourette's syndrome, hypertension, post-myocardial infarction, tachyarrhythmia, human immunodeficiency virus infection, AIDS, cell proliferation (cancer, psoriasis or hyperkeratosis), and drug abuse. (A) can also be used to treat peptic ulcer, glaucoma, asthma, renal disease, Parkinson's disease, congestive heart failure, migraine,
                                                                                                                                                                                                                                                                                                                                                                                                    27-MAR-1996;
25-JUN-1996;
20-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         isomerisation of prolyl peptide bonds in proteins and accelerates th folding of the protein. The inventors are claiming the PPIase-alpha. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-200 correct PF field.) (Updated on 25-MAR-2003 to correct PR field.) (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy; psychotic disorder; Huntington's disease; Parkinson's disease; post-myocardial infarction; HIV infection; cell proliferation disorder; peptic ulcer; glaucoma; asthma; renal disease; autoimmune disease; antibacterial; betal-adrenergic receptor.
                                                                                                                                                                                                                                    schizophrenia,
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(SEEM/)
(GEOR/)
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96US-00670119.
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The invention relates to identifying agents which may be potentially proceed test agent on the complexes of BAK and/or M11L proteins. The methods are useful for identifying agents which may be potentially pro-apoptotic or drug discovery system. The methods are conducting a target or drug discovery system. The methods are useful for identifying M11L-interacting polypeptides, and in conducting a target or drug discovery system. The methods are useful for identifying agents capable of inhibiting M11L activity or which can mimic the activity of M11L by inhibiting the activity of BAK and which are therefore anti-apoptotic agents. Agents identified by the method as involved in regulation of apoptosis may be used in the development of therapeutic agents and methods, and drug screening assays, and in the cactivity of cancer cells to chemotherapeutic treatment. Therapeutic applications of apoptosis manipulation include treatment of acute and chronic neurodegenerative diseases, e.g. stroke, Alzheimer's or funtington's disease by drugs, and sensitization of cancer cells for drug (radiation-induced apoptosis by modulation of survival signals and viral transfer of apoptosis promoting genes. The present sequence represents a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying agents which may apoptotic for treating acute comprises determining the eff and/or M11L proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
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                                  The present sequence represents a specifically claimed peptide sequence, peptide 113-F4.5.12, where at least one amino acid is in the D-CC configuration. The peptide preferably is modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The peptide, based on the naturally occurring histidine-rich human CC histatins, have anti-fungal and anti-bacterial activity and are useful in compositions for the treatment of oral, vaginal, urethral, mucosal, crespiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against local and systemic Candida albicans infection, against oral bacterial diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas gingivalis) and against Clostridium histolyticum. The peptide has cartivity, particularly on a weight basin, compared to the longer, naturally occurring histatins. Peptides containing D-residues are also more resistant to degradation than L-amino acid versions
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RESULT 11
AAW19464
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The present sequence represents a specifically claimed example of an anti-fungal and anti-bacterial D-amino acid histatin-based peptide 113-CC F4.5.12, where at least one amino acid in the peptide is in the D-CC configuration. The peptide is preferably modified by an acetyl or carbamyl addition at the N-terminus and/or amidation at the C-terminus. The novel D-amino acid-containing peptide, based on the naturally occurring histidine-rich human histatins, have anti-fungal and anti-CC bacterial activity and are useful in compositions for the treatment of coral, vaginal, urethral, mucosal, respiratory, skin, ear and ophthalmic fungal or bacterial infections. It is particularly effective against CC diseases such as caries and periodontitis (by inhibiting e.g. Streptococcus mutans and Porphyromonas gingivalis) and against CC clostridium histolyticum. The D-amino acid-containing peptide has superior anti-fungal (especially anti-candidal) and anti-bacterial activity, particularly on a weight basis, compared to the natural L-camino acid forms of histatins and histatin-based peptides. The presence of D-residues also makes the peptides more resistant to degradation than
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                                                                                                                                                                                                                                                                                                                                                               Anti-fungal and anti-bacterial D-amino acid histatins and histatin-based peptide(s) - useful esp for treating Candidal infections, periodontitis
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                                                                                                                                                                                                                                                                                                                  Claim 2; Fig 1; 63pp; English
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configuration"
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Pred. No. 3.6e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Roberts FD,
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RESULT 12
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                                                                                                                                                                               CC absent; R2 and R20 are Ser or absent; R3, R18, R19 and R21 are His or CC absent; R4 is Ala; R5, R6, R9, R12, R13, are Lys, Gln, Arg or another CC basic amino acid; R7, R8 and R15 are His, Phe, Tyr, Leu, another CC hydrophobic amino acid, or is absent; R10 is Tyr; R16 is Glu or absent; CC R17 is Lys or absent; R22 is Arg or absent; R23 is Gly or absent; where CC amino acid sequence. (A) can be used to treat fungal or bacterial CC infection in a mammal. The peptides are particularly used to treat CC vaginal, urethral, mucosal, respiratory, skin, ear, oral, or ophthalmic CC infections. The infections that can be treated include those caused by CC viscosus, Bacteriodes forsythus, Bacteriodes gracilis, Bacteriodes CC viscosus, Bacteriodes forsythus, Bacteriodes gracilis, Bacteriodes cohracea, Campylobacter sputorum, Capnocytophaga gingivalis, Capnocytophaga corrodens, Eubacterium nodatum, Fusobacterium nucleatum, Fusobacterium fusobacterium nucleatum, Fusobac
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Best Local S
Matches 5
   peridonticum, Peptostreptococcus nicros, Porphyromonas endodontalis, Porphyromonas gingivalis, Prevotella intermedia, Prevotella nigrescens, Propionibacterium acnes, Pseudomonas aeruginosa, and those caused by various Staphylococcal and Streptococcal species. AAY25140-Y25149 are peptides used in the description of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel substantially pure antibacterial and antifungal peptides (A) containing between 13 and 20 amino acids, where the peptide has the amino acid sequence: R1-R2-R3-R4-R5-R6-R7-R8- R9-R10-R11-R12-R13-R14-R15-R16-R17-R18-R19-R20-R21-R22-R23; where: R1 is Asp or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 8; 31pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (PERI-) PERIODONTIX INC (UYBO-) UNIV BOSTON.
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                                                                                                                                                                   Porphyromonas endodontalis,
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of a histatin, a tradgment of histatin, or a histatin, related peptide. The competide that is administered contains between 8 and 20 amino acids and CC has the sequence: X1-X2-X3-X4-X5-X6-X7-X8-X9-X10-X11-X12-X13-X14-X15-X16-X17-X18-X19-X20-X21-X22-X23 where X1 is Asp or absent; X2 is Ser or CC absent; X3 is His or absent; X4 is Ala, His, Leu or absent; X5 is Lys, CC Gln, Arg, Orn, or another basic amino acid; X6 is Arg, Gln, Lys, or another basic amino acid; X7 is His, Phe, Tyr Leu, or another hydrophobic camino acid; X8 is His, Phe, Tyr Leu, or another hydrophobic composition, or another hydrophobic amino acid; X12 is Lys, Gln, Arg, Corn, or another hydrophobic amino acid; X12 is Lys, Gln, Arg, Corn, or another hydrophobic amino acid; X12 is Lys, Gln, Arg, Corn, or another hydrophobic amino acid; X13 is Lys, Gln, Arg, Corn, or another hydrophobic amino acid; X13 is Lys, Gln, Arg, Corn, another hydrophobic amino acid, or absent; X16 is Glu or absent; X17 is Lys or absent; X18 is His or absent; X19 is His or absent; X17 is Lys or absent; X18 is His or absent; X19 is His or absent; X10 is Ser Corn, another hydrophobic amino acid, or absent; X16 is Glu or absent; X17 is Lys or absent; X17 is Lys or absent; X18 is His or absent; X19 is His or absent; X10 is Ser Corn, another hydrophobic amino acid, or absent; X16 is Glu or absent; X17 is Lys or absent; X17 is Lys or absent; X10 is Glu or absent; X10 is His or 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to treating cystic fibrosis via the administration of a histatin, a fragment of histatin, or a histatin-related peptide. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 32; Page 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-465852/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Spacciapoli P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200040204-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antibiotic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating cystic fibrosis and combating Pseudomonas and other pulmonary infections in a mammal comprises administering histatin or a fragment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-JUL-2000
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5; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rothstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   peptide; cystic fibrosis; Pseudomonas aeruginosa;
tance; pulmonary infection; antibacterial; antibiotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide, P-113-F4.5.12
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Pred. No.
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RESULT 14
AAW33942
ID AAW33
XX AAW33
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This sequence is a betal-adrenergic receptor antagonist, and is an CC antagonist of the invention. The antagonists (A) are for inhibiting the function of a prokaryotic or eukaryotic integral membrane protein (IMP) having at least one transmembrane domain (TMD), comprises a peptide concluding at least 4 consecutive amino acids (aa) from the sequence of the TMD. (A) are used to treat or prevent disorders in mammals that con the treat or prevent disorders in mammals that conditions that can be treated are schizophrenia, psychotic disorders, conditions that can be treated are schizophrenia, psychotic disorders, infarction, tachyarrhythmia, human immunodeficiency virus infection, call proliferation (cancer, psoriasis or hyperteratosis), and drug abuse: (A) can also be used to treat peptic ulcer, glaucoma, asthma, can be used to treat peptic ulcer, glaucoma, asthma, disease, graft rejection etc. (A) can be used as diuretics and consenses, graft rejection etc. (A) can be used as diuretics and consenses, graft rejection etc. (A) can be used as diuretics and consenses, when attached to a marker, for tissue imaging to localise/quantify specific receptors. The transgenic animals are used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
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25-JUN-1996;
20-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            therapy; psychotic disorder; Huntington's disease; post-myocardial infarction; HIV infection; cell propertic ulcer; glaucoma; asthma; renal disease; autoantibacterial; betal-adrenergic receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NGGY/)
(SEEM/)
(GEOR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 17; Page 94; 127pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Inhibitor of integral membrane protein - used to treat schizophrenia, hypertension, viral infection, cancer et
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ng GYK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9735881-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ODOW/)
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SEEMAN
GEORGE
O'DOWD
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96US-00670119.
96US-0024240P.
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62.5%;
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Pred. No.
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RESULT 15
AAW68100
ID AAW68
XX AAW68
XX AAW68
XX AAW68
AC AAW68
CDT 27-00
XX Chla
XX Fey
XX Fey
YN WO9
XX Chl;
XX I6
XX I7
PR 12
XX YF 16
XX YF 17
PR 12
XX YF 16
XX YF 17
YF 
RESULT 16
AAW16616
ID AAW16
XX
AC AAW16
XX
DT 19-DE
XX
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Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chlamydomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chlamydomonas reinhardtii RB47 binding protein peptide fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RB47; activator; mRNA binding regulated production; protein
                           19-DEC-1997
                                                                         AAW16616;
                                                                                                                    AAW16616 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence is that of a fragment of the RB47 mRNA binding protein was used in the isolation of the coding region of the complete gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 58; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plant cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated RB47 and RB60 genes - used for, e.g. regulated expression desired recombinant molecule in prokaryote and eukaryote, particularly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-427562/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mayfield
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JAN-1997;
12-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9831823-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW68100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SCRI ) SCRIPPS RES INST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ω
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5; Conserv
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~•
                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                           GFGFINFK 8
                                                                                                                                                                                                                                                                                         GFGYKNSK 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GYANSAF 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GYKNSKF 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                     AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reinhardtii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0035955P.
97US-0069400P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US000840
                                                                                                                  peptide; 11
                                                                                                                                                                                                                                                                                                                                                              52.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       52.9%;
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                                                                                                                                                                                                                                                                                                                                      o
••
                                                                                                                                                                                                                                                                                                                                      Score 27; DB 2;
Pred. No. 4.3e+02;
1; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; psbA; chloroplast; recombinant;
synthesis; plant propagation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; DB 2;
J. 4.3e+02;
2;
                                                                                                                                                                                                                                                                                                                                                                                    Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    indels
                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein which
                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       O ff
                                                                                                                                                                                                                                                                                                                                      0;
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RESULT 17
AAO22420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A novel interaction site has been discovered on phosphoinositide 3 (PI-3) kinase, or a homologue or analogue. The interaction site modulates the activity of PI-3 kinase when exposed to a modulator, and has a molecular shape adapted to interact with at least a part of the modulator so as to modulate PI-3 kinase activity. The present sequence represents a cc conserved motif (resembling K(X)nKXKK where n=3-7) in PI-3 kinase cc piloalpha, that was found to bind phosphatidylinositol in gelsolin and so might constitute a binding site for the 4,5-phosphates of the lipid. The activity of PI-3 kinase can be regulated by altering, e.g. substituting a cc site may be used to identify or design novel ligands which regulate the activity of PI-3 kinase by generating a molecular model of the wortmannin in interact with at least part of the site and optionally contacting the putative ligand with PI-3 kinase and monitoring PI-3 kinase activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                              Cytostatic; antiinflammatory; antibacterial; matrix metalloproteinase; protease cleavage site motif; enzyme; proteolytic enzyme; pathogenesis; B. anthracis; pla; YopJ protease; Yersinia; smallpox HIL metalloprotease; cathepsin family protease; TACE; calpain; caspase; BACE; beta-secretase; tumour necrosis factor-alpha converting enzyme; blood clotting cascade; beta-site amyloid precursor protein cleaving enzyme; presenilin; furin; membrane-type serine protease; proprotein convertase; protease inhibitor; proteasome; pathogenic infection; cancer; inflammatory disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 32; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Phospho:inositide 3 kinase wortmannin interaction site - to identify and design ligands which regulate phospho:inositide 3 kinase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wymann MP, 1
Zvelebil MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                        Protease cleavage site related peptide SEQ ID No 86.
                                                                                                                                                                                                                           11-OCT-2002
                                                                                                                                                                                                                                                                                            AAO22420 standard; peptide; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-259013/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Phosphoinositide 3 kinase; PI-3 kinase; wortmannin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Phosphoinositide-3 kinase pll0alpha conserved motif.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (LUDW-) LUDWIG INST CANCER RES
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                                                                                                                                                                                                                                                                                                                                                                                                                   FGYKNSK 8
                                                                                                                                                                                                                                                                                                                                                                                    FGYKRER 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bulgarelli-Vela G,
, Waterfield MD;
                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       51.0%;
57.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>'</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 26; DB 2;
Pred. No. 5.1e+02;
                                                                                                                                                                                                                                                                                               B
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vanhaesebroeck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 11;
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RESULT 18
AAM97164
ID AAM97
XX
AC AAM97
XX
AC AAM97
XX
AC Human
DT 24-JA
DT 24-JA
DT UMMAN
CAMMENT
CAM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  motif for a cleavage site of a protease. The method comprises contacting the protease with a peptide library containing degenerate residues which allow for cleavage of a substrate by the protease, allowing the protease to cleave peptides within the degenerate peptide library having a cleavage site for the protease to form a population of cleaved peptides, and determining an amino acid sequence motif for the cleavage site of the protease. The protease of the method is a matrix metalloproteinase, or a protease. The protease of the protease is a lethal factor of B. anthracis, pla and YopJ proteases of Yersinia, and the smallpox HIL metalloprotease. The protease of the invention is selected from a protease of pathogenic organisms, cathepsin cappains, caspases, beta-site amyloid precursor protein cleaving enzyme (BACE, beta-secretase), presentlins, membrane-type serine proteases, furin and other proprotein convertases, proteaseme components and proteases affecting the blood clotting cascade. The protease inhibitors of the invention are useful to treat diseases, including pathogenic components a peptide sequence relating to the protease cleavage site motifs of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                        Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cano amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f;
                                                                                                                                                                                                                                                                                                                           Human peptide #439 encoded by a SNP oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 12
                                                      nervous system
                                                                                           multifactorial
                                                                                                                                                                                                                                                                                                                                                                                              24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAM97164;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAM97164 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 8; Page 52; 126pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-NOV-2001; 2001WO-US046777
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                                                                                                                         interleukin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BETH-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                determination of oriented peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BETH ISRAEL DEACONESS MEDICAL CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FGYKRER 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FGYKNSK 8
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                                                                                                                                                     related
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                 elated protein; cytochrome; kinesin; cytokine; interferon; G-protein coupled receptor; thioesterase; inflammation; al disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0246815P.
                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 51.0%;
57.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protease cleavage site motifs using a mixture-library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a method for determining an amino aci
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Pred. No. 5.5e
1; Mismatches
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5.5e+02;
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                                                                                                                                                                                                                                       cancer;
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Homo sapiens.

27-SEP-2001

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CC encoding polymorphic variants of proteins related to amylases, amyloid
CC proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin,
CC polymerase, oncogenes, histones, kinases, colony stimulating factors,
CC complement related proteins, cytochromes, kinesins, cytokines,
CC interferons, interleukins, G-protein coupled receptors and thioesterases.
CC The present sequence is a peptide encoded by one such oligonucleotide.
CC The oligonucleotides and the peptides encoded by them may be used in the
CC prevention, diagnosis and treatment of diseases associated with
CC inappropriate expression of the proteins listed above. Disorders that may
CC be prevented, diagnosed and/or treated include multifactorial diseases
CC with a genetic component, such as autoimmune diseases (e.g. rheumatoid
CC arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus
and Grave's disease), inflammation, cancer (e.g. cancers of the bladder,
CC brain, breast, colon and kidney, leukaemia), diseases of the nervous
CC system and an infection of pathogenic organisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                              Glutamic acid; glutamine; vaccine; major histocompatibility complex; MHC; immunomodulator; antiallergic; endocrine; neuroprotectant; virucidal; bactericidal; antiparasitic; fungicidal; cytostatic; medicine; pharmaceutical; immune disorder; immune deficiency; autoimmune; hypersensitivity; allergy; graft rejection; infection; hormonal disorder; central nervous system disease; cancer; melanoma; anti-melanoma vaccine; human immunodeficiency virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic nucleic acids encoding oncogenes and histones, useful for autoimmune diseases and infections.
                                                                                                                                                                                                                                 Vaccine related MHC ligand peptide SEQ ID NO:642.
                                                                                                                                                                                                                                                                     07-DEC-2001
                                                                                                                                                                                                                                                                                                      AAM99539;
                                                                                                                                                                                                                                                                                                                                     AAM99539 standard; peptide; 14 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 14 AA;
                                                                Clostridium tetani.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUL-2001
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27-DEC-2000; 2000US-00173419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GFGYKNSKF 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       . Similarity 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                         GAEYDNNKF 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 3764; 4143pp; English.
                                                                                                                                                                                                                                                                     (first
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                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           51.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 26; DB
Pred. No. 6.5e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e.g. amylases, diagnosing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 4;
6.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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, e.g. cancer,
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RESULT 20
ADB71196
ID ADB71
XX
AC ADB71
XX
AC ADB71
XX
DT 04-DE
XX
Varia
KW varia
KW activ
KW coagu
XX
Coagu
XX
Homo
XX
Homo
XX
PD 12-S1
XX
PF 28-F1
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          acid (II). Also described are: (a) a pharmaceutical composition

CC containing at least one (I); (b) a vaccine containing at least one (I)

CC where this is a major histocompatibility complex (MHC) ligand (Ia); (c) a

CC (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process

CC for preparing (I). (I) has immunomodulator, endocrine, antiallergic,

CC neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and

CC cytostatic activities. (I) are useful, in human or veterinary medicine,

CC immune deficiency, autoimmune states, hypersensitivity, allergy, graft

CC rejection, infection, hormonal disorders and central nervous system

CC diseases), also, where (I) is a MHC ligand (Ia), in vaccines for

CC treatment or prevention of: (i) viral, bacterial, parasitic or fungal

CC melanoma vaccines. (I) are also useful for in vitro diagnosis of diseases

CC melanoma vaccines. (I) are also useful for in vitro diagnosis of diseases

CC melanoma be used in pharmaceutical compounds from the present peptides

CC which can be used in pharmaceutical compounds from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 5
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                                                                                                                                                                              variant blood coagulation component; anticoagulant; protein activated protein C; blood; thrombolytic; anticoagulant; coagulation disorder; thrombosis; mutant; variant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a pharmaceutical compound (I) that contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue the form of an addition salt with a strong, physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 9; Page 140; 149pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-611470/70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-MAR-2001; 2001WO-FR000872
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                    28-FEB-2003; 2003WO-SE000331
                                                                                           WO2003073980-A2
                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                     Human modified protein C peptide SEQ ID NO:42
                                                                                                                                                                                                                                                                                           04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                               ADB71196 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 14 AA;
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                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                               peptide; 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         51.0%; Score 26; DB 4; Lo
100.0%; Pred. No. 6.5e+02;
live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      containing N-terminal glutamic acid or anti-melanoma vaccines, is an addition
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Glutamic acid;

07-DEC-2001

(first entry)

AAM99439;

AAM99439 standard; peptide; 9 AA

Vaccine related MHC ligand peptide SEQ ID NO:542.

immunomodulator; antiallergic; endocrine; neuroprotectant; virus bactericidal; antiparasitic; fungicidal; cytostatic; medicine; pharmaceutical; immune disorder; immune deficiency; autoimmune; hypersensitivity; allergy; graft rejection; infection; hormonal central nervous system disease; cancer; melanoma; anti-melanoma human immunodeficiency virus.

disorder;

glutamine; vaccine; major histocompatibility complex;
r; antiallergic; endocrine; neuroprotectant; virucidal;

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ω

1 GFGYKNSK 8

Matches Query Match Best Local

4.

Conservative

Similarity

51.0%; 50.0%;

Score 26; I Pred. No. 6. Mismatches

DB 7; 6.5e+02;

Length 14;

<u>ب</u>

0

Gaps

0

Sequence 14 AA;

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which is substantially homologous in amino acid sequence to a wild-type or anticoagulant protein in the protein C (APC). (I) exhibits enhanced canticoagulant activity compared to the wild-type blood component, and component activity compared to the wild-type blood component, and component adiabation in a region corresponding to the serine-protease (SP) domain of the wild-type protein. Also described: (I) a DNA segment comprising a nucleotide sequence encoding (I); (2) a DNA segment (C); (4) producing the DNA segment coding for the variant blood coagulation component by introducing mutations into the containing the DNA segment of (1); (3) a mixroorganism or (preferably an) animal host cell stably transfected (I) from a culture of the transfected host cell; (6) a diagnostic test system, especially a kit, for assaying components participating in the protein C containing (I). (I) has thrombolytic and anticoagulant activities. The medicament for treating or preventing coagulation disorders, such as thrombosis. The medicament is especially suitable for treating a patient of the protein C anticoagulant system of blood, composition of a composition in the protein C anticoagulant system is useful in the manufacture of a containing in the protein C anticoagulant for assaying components participating a patient coagulation disorders, such as thrombosis. The medicament is especially suitable for treating a patient coagulation disorders, such as thrombosis in the protein C anticoagulant activity compared to the wild-type protein. The present sequence represents a human modified protein C peptide, which is given in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New variant blood coagulation component substantially homologous to protein {\tt C} or activated protein {\tt C} useful for preparing a composition treating or preventing coagulation disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a variant blood coagulation component which is substantially homologous in amino and a component
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAR-2002; 2002US-0360181P.
06-AUG-2002; 2002US-0401042P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 46; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TACT-) TAC THROMBOSIS & COAGULATION AB
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RESULT 22
AAM99577
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9 AA;
             Glutamic acid; glutamine; vaccine; major histocompatibility compimmunomodulator; antiallergic; endocrine; neuroprotectant; virus bactericidal; antiparasitic; fungicidal; cytostatic; medicine; pharmaceutical; immune disorder; immune deficiency; autoimmune; hypersensitivity; allergy; graft rejection; infection; hormonal central nervous system disease; cancer; melanoma; anti-melanoma human immunodeficiency virus.
                                                                                                                                                             Vaccine related MHC ligand
                                                                                                                                                                                                07-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stabilized pharmaceutical containing N-terminal glutamic acid or glutamine, useful e.g. in anti-melanoma vaccines, is an addition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                     AAM99577 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-MAR-2001; 2001WO-FR000872
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                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                     peptide; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   49.0%;
                                                                                                                                                           peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 25;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beck A,
                                                                                                                                                             SEQ ID NO:680.
                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4; Lu., 1.4e+06; 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                    virucidal;
                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                               disorder; vaccine;
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RESULT 23
AAW28520
ID AAW28
XX
AC AAW28
XX
DT 11-JA
XX
DE Human
XX
Biose
KW Biose
KW Herpe
XX
Herpe
XX
Synth
XX
PN AT950
XX
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immune deficiency, autoimmune states, hypersensitivity, allergy, graft rejection, infection, hormonal disorders and central nervous system diseases), also, where (I) is a MHC ligand (Ia), in vaccines for treatment or prevention of: (i) viral, bacterial, parasitic or fungal infections; or (ii) of cancers. A particular application is in antimelanoma vaccines. (I) are also useful for in vitro diagnosis of diseases associated with interactions between MHC and (I), e.g. melanoma and human immunodeficiency virus infection. AAM98898 to AAM99592 represent peptides which can be used in pharmaceutical compounds from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a pharmaceutical compound (I) that contains an N-terminal glutamic acid (Glu) or glutamine (Gln) residue in the form of an addition salt with a strong; physiologically acceptable acid (II). Also described are: (a) a pharmaceutical composition containing at least one (I); (b) a vaccine containing at least one (I) where this is a major histocompatibility complex (MHC) ligand (Ia); (c) method for in vitro diagnosis of diseases associated with the presence (Ia); (d) a kit for method (c) that includes a (Ia); and (e) a process for preparing (I). (I) has immunomodulator, endocrine, antiallergic, neuroprotectant, virucidal, bactericidal, antiparasitic, fungicidal and cytostatic activities. (I) are useful, in human or veterinary medicine, in pharmaceutical compositions (for treating immune disorders, exactivity automorphic parasity and compositions of the compositions o
                                                                                                                                                 Biosensor; capture; ligand;
Herpes virus.
                                                                                                                                                                                                                            Human Herpes VI sequence
                                                                                                                                                                                                                                                                              11-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                AAW28520 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 9 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stabilized pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Klinguer-Hamour C,
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                                                                                                                                                                                                                                                                                                                                                                            peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                49.0%; but
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Corvaia N,
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                                                                                                                                                                          ion; electron; channel; current;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 25;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beck
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. 1.4e+06;
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O D

AT9501943-A Synthetic.

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RESULT 24
AAW63824
ID AAW63
XX
AC AAW62
XX
DT 28-SJ
XX
DE Huma
XX
Inte
KW Inte
KW Inmu
KW Monn
XX
OS Hom
XX
PP 23
PP 23
XX
PP 21
PP 26
PR 27
XX
PP XX
PP T 1
PT 1
PT T
XX
PS
X
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                           26-DEC-1996;
21-FEB-1997;
06-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 12 describes the prodn. of a Herpes virus biosensor where the capture ligands (oligonuclectides) are based on the sequences given in AAW28515-W28526. The biosensor is useful for determining DNA or RNA concentrations by a method comprising measuring changes in ion and/or electron current induced by hybridisation in the channels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SCHA/)
(PITT/)
                                                                                                                          New interleukin-18 receptor protein - used to inhibit interleukin-18, treat autoimmune disease and as immunosuppressant to treat e.g. graft rejection, insulin-resistant diabetes and ulcerative colitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bio:sensor for nucleic acid determination - with capture ligands immobilised in ion channels in membrane.
    This sequence represents the human monoc heavy chain complementarity determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunosuppressant; inhibitor; receptor protein; detection; heavy ch monoclonal antibody; Mab; complementarity determining region; CDR3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Mab #117-10C heavy chain CDR3 peptide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 12; Page 9; 17pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-NOV-1995;
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                                                                                                                                                                                                                                   WPI; 1998-335317/30.
                                                                                                                                                                                                                                                                                  Torigoe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW63824;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW63824 standard;
                                                                                                                                                                                                                                                                                                                                       (HAYB ) HAYASHIBARA SEIBUTSU KAGAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Interleukin-18:
                                                                               14;
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                                                                                                                                                                                                                                                                                Ushio S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                         96JP-00356426.
97JP-00052526.
97JP-00163490.
                                                                               20;
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                                                                                                                                                                                                                                                                                                                                                                                      97JP-00215490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97EP-00310555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IL-18; human; treatment; autoimmune disease; Mab;
nt; inhibitor; receptor protein; detection; heavy chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptide; 10
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                                                                                                                                                                                                                                                                                  Kunikata T,
                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 25; DB
Pred. No. 7e+0
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ۲.
        monoclonal antibody (Mab) #117-10C ining region, CDR3 which is used in
                                                                                                                                                                                                                                                                                     Kurimoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2;
7e+02;
      region,
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RESULT 25
AAB46262
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diseases, Goodpasture's syndrome, primary biliary cirrhosis, sympathetic ophthalmitis, hyperthyroidism, juvenile onset type diabetes, Sjogren syndrome, autoimmune hepatitis, autoimmune haemolytic anaemia, myastheni gravis, systemic scleroderma, systemic lupus erythematosus, polyleptic cold haemoglobinuria, polymyositis, periarteritis nodosa, multiple sclerosis, Addison's disease, purpura haemorrhagica, Basedow's disease, leukopaenia, Behcet's disease, climacterium praecox, rheumatoid arthritis, rheumatopyra, chronic thyroiditis, Hodgkin's disease, allergy and septic shock resulting from production or administration of excessive gamma interferon (IFN-gamma)
This invention describes novel T cell epitopes (A) of the human papilloma virus II protein. The invention also describes (a) compounds (I) containing an (A) provided it is not a natural II protein nor an exclusively N- or C-terminal deletion variant of natural II; (b) nucleic acid (II) that encodes (A) or (I); (C) vector, particularly an expression vector, that contains (II); (d) cells that contain, and preferably present, (A); (e) complex of (A) or (I) with at least one additional molecule (III); in vitro detection of T cell activation by a compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   method involved in neutralising IL-18 or to treat autoimmune diseases or as an immunosuppressant using anti-IL-18 antibodies which can inhibit IL-18. Such antibodies can also be used to detect the IL-18 receptor protein (labelled with an enzyme or a radioactive or fluorescent substance). The protein is used to treat e.g. graft rejection, pernicious anaemia, atrophic gastritis, insulin-resistant diabetes, Wegener granulomatosis, discoold lupus erythematosis, ulcerative colitis, cold-agglutinin-relating discoold lupus erythematosis, ulcerative colitis, cold-agglutinin-relating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cytotoxic T cell; epitope; vaccine; tumor; protective
                                                                                                                                                                                                                                                                                                                                                                                                 01-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                   07-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human papillomavirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HPV type 16 cytotoxic T-cell epitope SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB46262;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB46262 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10 AA;
                                                                                                                                                                                          detecting
                                                                                                                                                                                                         New T cell
                                                                                                                                                                                                                                              WPI; 2001-062580/08
                                                                                                                                                                                                                                                                            Schaefer K,
                                                                                                                                                                                                                                                                                                               (MEDI-)
                                                                                                                                                                                                                                                                                                                                                                01-JUN-1999;
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                                                                                                                                                         Example 4; Page 18; 26pp; German.
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DKFZ DEUT KREBSFORSCHUNGSZENTRUM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 4; Conserv
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                                                                                                                                                                                       l epitopes from or inducing an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                             Faath S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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100.0%; Pr
                                                                                                                                                                                                                                                                              Jochmus I,
                                                                                                                                                                                          the papilloma virus L1 protein, us immune response, e.g. in vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        L1 protein; antiviral; antitumor; antigen; immune response.
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0; Mismatches
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                                                                                                                                                                                                                                                                                Nieland J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
7e+02;
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RESULT 26
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PF 18-JI
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PF 19-JI
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PF 19-JI
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PF 10-JI
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PF 10-SE
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PF 10
               The present invention describes an isolated salivary polypeptide (I), of Phlebotomus papatasi. Also described: (I) an isolated nucleic acid (II) encoding (I); (2) an isolated fragment (III) of (II), where the fragment encodes the polypeptide fragment specific for a polypeptide such as Phlebotomus papatasi salivary polypeptide (PpSP) 12, PpSP14, PpSP15, PpSP30, PpSP32, PpSP36, PpSP42 or PpSP44; (3) an isolated antigenic or immunogenic fragment (IV) of (I); (4) a nucleic acid (V) that hybridises under stringent conditions to (II); (5) a vector (VI) comprising (II) or (III); and (6) a composition (VII) comprising (I) or its fragment, (IV) or (VI), and a pharmaceutically acceptable carrier. (I) has protozoacide and immunostimulant activities, and can be used in vaccines. (VII) is useful for preventing Leishmaniasis in a subject. (II) is also useful for preventing Leishmaniasis in a subject. (VII) is useful as a vaccine. ACC79987 to ACC79999 and ABR91176 to ABR92046 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 containing a system for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 10; Page 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR91208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated salivary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                papatasi; salivary polypeptide; antigenic;
e; immunostimulant; vaccine; immune response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vary polypeptide of Phlebotomus papatasi, response in a subject or for preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kamhawi S,
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Pred. No. 7e+0
1; Mismatches
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7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , useful for
Leishmaniasis
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RESULT 28
AAB81463
ID AAB81
XX
AC AAB81

standard;

peptide; 12

AAB81463; AAB81463

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RESULT 27
AAB81464
ID AAB81
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Matches 4
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Best Local S
Matches 3
                                                                                                                                                                                                                                    The present sequence is part of a template-fixed, beta-hairpin loop peptidomimetic which may be manufactured by the process disclosed in this invention. The process is based on a mixed solid and solution phase synthetic strategy. The method is useful for determining key amino acids and motifs important for binding large surface and flat protein interfaces in their sequential and/or spatial arrangement. This information can ultimately be used for the design of small peptidomimetic drug candidates. The peptidomimetics may be used to probe large surface protein-protein interactions and to find protein targets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Method
useful
                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Robinson JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptidomimetic; beta-hairpin loop mimetic; binding motif identification; drug design; protein target identification.
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  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for manufacturing template-fixed beta-hairpin loop mimetics, for designing small peptidomimetic drug candidates, involves based on mixed solid and solution phase synthetic strategy.
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3; Conserv
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                                                                                              Similarity 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page
YGVKUSEW
                                                                                                                                                                                                 12
                                                 FGYKNSKF 9
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                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-EP006369
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  11
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                                                                                                Score 25; DB Pred. No. 8.4e 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 25;
Pred. No.
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7e+02;
                                                                                                                         8.4e+02;
                                                                                                                                                    DB 4;
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                                                                                                                                                Length 12,
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                                                                                              Gaps
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Method for manufacturing template-fixed beta-hairpin loop mimetics, useful for designing small peptidomimetic drug candidates, involves process based on mixed solid and solution phase synthetic strategy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic peptide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptidomimetic; beta-hairpin loop mimetic; binding motif identification; drug design; protein target identification.
                                                                             Synthetic
                                                                                                                    Fusogenic; nuclear localisation signal; NLS; encapsulation; lipogene; liposome; micelle; karyophilic; cytostatic; antitumour; solid tumour; peptide-lipid-polynucleotide complex; neoplastic disease; gene therap;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 7; Page 77; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-273332/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Robinson JA,
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   08-JUN-2001; 2001WO-US018657
                                                                                                                                                                         Simple nuclear localisation
                                                                                                                                                                                                   18-APR-2002
                                                                                                                                                                                                                                                     ABB74314 standard; peptide; 12 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 12 AA;
                                                     WO200193836-A2
                                                                                                         breast
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                                                                                                         carcinoma;
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                                                                                                                                                                                                                                                                                                                        YGVKONSEW 11
                                                                                                                                                                                                                                                                                                                                                 FGYKNSKF 9
                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                        prostate
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                                                                                                                                                                                                                                                                                                                                                                                       49.0%;
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                                                                                                                                                                        signal peptide SEQ ID NO:78
                                                                                                                                                                                                                                                                                                                                                                        Score 25; DB 4; LC
Pred. No. 8.4e+02;
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                                                                                                                      gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for producing micelles with centrapped therapeutic agents. The method comprises: (1) combining confidence of the negatively charged atoms are neutralised by positive charges on lipid molecules to form an electrostatic micelle complex in 20-80 % continuing the micelle complex of (a) with fuscioning the micelle complex of (a) with fuscioning confidence in a 0.0-0.3 ratio, therefore producing micelles with entrapped therapeutic agents. Also described is a method confidence in the micelle complex of (a) with fuscioning confidence in the micelle complex in the respection of the micelle. ABB74256 to ABB74858 represent specifically claimed conclear localisation signal (NLS) peptides for use in the method as the fuscionic karyophilic peptides. The micelles produced can have cytostatic and antitumour activities. The peptide-lipid-polynucleotide complexes produced are useful for inhibiting the progression of neoplastic diseases. The invention relates to the field of gene therapy and is complexes suitable for delivery of polynucleotides. The encapsulated complexes suitable for delivery of polynucleotides. The encapsulated concluding but not limited to breast carcinoma or prostate carcinoma. ABB74255 are used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12 AA;
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                                                                                                                                                                                             Leucoanthocyanidin reductase; LAR; reductase-epimerase-dehydrogenase; RED; proanthocyanidin; pasture quality; legume; bloat-safe forage crop; disease resistance; pest resistance; enzyme.
                                                                                                                                                                                                                                                             Desmodium uncinatum leucoanthocyanidin reductase (LAR)
                                                                                                                                                                                                                                                                                                                                                                AAE28582 standard; peptide; 12
                                             Misc-difference
                                                                               Misc-difference
                                                                                                                                                               Desmodium uncinatum
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                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       liposome complexes.
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4; Conservative
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                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                             Location/Qualifiers
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                                                               label= Leu,
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Pred. No. 8.4e+02;
2; Mismatches 2;
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RESULT 31
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Major histocompatibility class II; MHC; binding; vaccine; MBP; autoimmune disease; myelin basic protein; HLA-DRB1*0401.
                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New LAR polypeptide from the RED protein superfamily, useful for regulating the biosynthesis and accumulation of proanthocyanidins in plants, and modifying the pasture quality of legumes.
Identifying MHC binding peptides - by ascertaining the characteristics a MHC molecule binding groove and determining conformation scores of
                                                                                WPI; 1999-105639/09
                                                                                                                                                                                                                          20-JUN-1997;
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                                                                                                                                                                            (ECLA-)
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Francki KT;
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                                                                                                                                                                                                                                                                                                                                                                                                              03-DEC-1998;
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                                                                                                                                                                                                                      WPI; 2000-423022/36
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100.0%;
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Diagnosing modulators

and treating obesity and impaired glucose tolerance using of daf-18 expression and/or activity.

The present sequence is conserved protein region

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Caenorhabditis

Page 171;

402pp; English

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The invention relates to a novel modified bacterial enzyme carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any non-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having alterations compared with the non-modified parochial enzyme. The alterations cause a reduction or an elimination of one or more of T-cell epitope sequences, which act in the parental enzyme as MHC class II binding ligands and stimulate T-cells. The modified CPG2 enzyme and the
                                                                                                                                                                                                                                                                                   WPI;
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                                                                                                                                                                                                               New modified bacterial enzyme substantially non-immunogenic CPG2, useful for inducing an i
                                                                                                                                                                                                                                                                                                                                                                                         29-NOV-2001; 2001EP-00128519.
25-JAN-2002; 2002EP-00001778.
13-SEP-2002; 2002EP-00020634.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand; immunostimulant; enzyme therapy; immune response; gene directed enzyme prodrug strategy; vaccine; enzyme; EC 3.
                                                                                                                                                                                                                                                                                                                       Hellendoorn
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                                                                                                                                                                               Claim
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                                                                                                                                                                              3; Page 13; 52pp; English.
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                                                                                                                                                                                                                                                                                                                       Baker M,
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                                                                                                                                                                                                              zyme carboxypeptidase G2 (CPG2) having enic or less immunogenic than any non-man immune response in a human host.
                                                                                                                                                                                                                                                                                                                          Williams
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Pred. No. 9.1e+02;
1; Mismatches 1
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The invention relates to a novel modified bacterial enzyme carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any non-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having alterations compared with the non-modified parochial enzyme. The alterations cause a reduction or an elimination of one or more of T-cell epitope sequences, which act in the parental enzyme as MHC class II binding ligands and stimulate T-cells. The modified CPG2 enzyme and the CPG2 proteins have immunostimulant activity and may be used in enzyme therapy. The modified CPG2 enzyme may be used to induce an immune response in a human host, or as a therapeutic entity such as the gene directed enzyme prodrug strategy. The peptide is useful for the
                                                                                                                                                                                                                                                                              New modified bacterial enzyme carboxypeptidase G2 (CPG2) having substantially non-immunogenic or less immunogenic than any non-modified CPG2, useful for inducing an immune response in a human host.
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                                                                                                                                                                                                                                                                                                                                                                                     Hellendoorn K,
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; 2002EP-00001778.
; 2002EP-00020634.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         #59 with potential human MHC class II binding activity.
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                                                                                                                                                                                                                                             52pp;
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                                                                                                                                                                                                                                                                                                                                                                                          Williams
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RESULT
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                                                                                 The invention relates to novel pseudopeptidic osteogenic growth polypeptide (OGP) analogues. The analogues of the invention may be used for stimulating the formation of osteoblastic or fibroblastic cells, enhancing bone formation in osteopenia-related pathological conditions, repairing fractures, healing wounds, grafting of intraosseous implants and reversing bone loss in osteoporosis and other conditions requiring enhanced bone cell formation. Furthermore, the analogues may have activities similar to or superior to that of the parent oligopeptides OGP(1-14) and OGP(10-14), as well as increased resistance to peptidase degradation, longer persistence in circulation or increased potency and bioavailability. The current sequence is that of the retro osteogenic growth polypeptide fragment of the invention.
                                                                                                                                                                                                                                                                                                     the
                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                     Bab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    osteogenic growth polypeptide; OGP analogue; osteoblastic; fibroblastic cell production; bone formation; osteopenia; fracture; wound; intraosseous implant; osteoporosis; OGP(1-14); OGP(10-14).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      manufacture of a modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo. This sequence represents a CPG2 enzyme peptide with potential human MHC class II binding activity of the invention.
                                                           Sequence
                                                                                                                                                                                                                                                             Example;
                                                                                                                                                                                                                                                                                    conditions requiring
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49.0%; brain 100.0%; Pre
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RESULT 37
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ID AAU25
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AAU25297

standard; peptide;

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                                                                                                                                  The invention relates to diagnosing, determining the stage or severity, or identifying the risk of a subject developing cancer (especially breast cancer), or monitoring the effect of therapy on a subject with cancer, comprising analysing a test sample using two-dimensional electrophoresis and detecting Breast Cancer-Associated Protein Isoforms (BPIS). The methods disclosed are used for the diagnosis and prognosis of breast cancer, for determining the severity of breast cancer, and monitoring cancer, for determining the severity of breast cancer, and monitoring the effect of therapy administered to a subject. Antibodies raised cancer the binding domain of a BPI, the binding domain of a BPI, a concleic acid encoding a BPI, or a nucleic acid that inhibits the function concleic acid encoding a BPI, or a nucleic acid that inhibits the function concleic acid encoding a BPI, or a nucleic acid that inhibits the function concleic acid encoding breast cancer. The methods use sensitive and compositions are more potent, specific, and has a more rapid effect with fewer side effects that other prior art methods. The present sequence is cancer side effects that other prior art methods. The present sequence is a tryptic digest peptide from a BPI of the invention
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                                                         Best
                                                                       Query Match
                                                                                                         Sequence 14 AA;
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24-MAR-2000; 2000GB-00007265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying proteins for clinical screening, diagnosis and prognosis breast cancer, comprises detecting Breast Cancer-Associated Protein Isoforms (BPIs) using two-dimensional electrophoresis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             breast
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                                                         Local
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ms (BPIs) using
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                                          Conservative
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tostatic; BPI; tryptic digest peptic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O'hare MJ,
                                                        49.0%;
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                                      Score 25; DB Pred. No. 9.8e 0; Mismatches
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                                                        DB 4;
9.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence represents a schizophrenia-associated protein isoform (SPI) These protein isoforms, e.g. SPI-206, SPI-238 and SPI-240 are detectable in cerebrospinal fluid, serum or plasma and are useful markers of schizophrenia. The sequences can be used for treatment and diagnosis of schizophrenia, screening, prognosis, monitoring the results of therapy, identifying patients most likely to respond to a particular therapy and identification of new targets for drug treatment. SPI DNA is useful as a nucleic acid probe to detect the presence of nucleic acids or SPIs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New schizophrenia associated protein isoforms and encoding nucleic acid molecules, useful for treatment, diagnosis and prognosis of schizophrenia and screening for potential drugs for treatment and new drug targets.
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28-DEC-2000; 2000US-00750395.
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                                                                                                               Schizophrenia; neuroleptic; diagnostic; neuropsychiatric disorder; neurological disorder; neuropathy.
                                                                                                                                                     Schizophrenia-associated isoform peptide #526
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14
               23-FEB-2001; 2001WO-GB000783.
                                        30-AUG-2001
                                                                WO200163293-A2
                                                                                        Homo sapiens
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                                                                                                                                                                                                      AAU15641;
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80.0%;
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9.8e+02;
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The invention relates to methods and compositions for screening, continuous and prognosis of Schizophrenia. The method involves detecting the presence of Schizophrenia (SCH) Associated Features (SFS) and SCH Associated Protein Isoforms (SPIS) in samples, e.g. by electrophoresis, continuous your hybridisation assay, for diagnosing and monitoring SCH, studying the effectiveness of treatments and for identifying potential therapeutic agents. The method is used for (1) screening or diagnosis of SCH and the relative abundance of at least 1 chosen feature correlates with the presence or absence of at least 1 chosen feature which correlates with the severity of SCH. The expression and activity of the SFS, SPIS and relative abundance of identify and produce potential therapeutic agents for the progress of the disorder and the effectiveness of treatment and as targets to identify and produce potential therapeutic agents for the treatment of SCH. The paucity of detectable neuralgic defects distinguishes neuropsychiatric disorders such as SCH from neurological disorders, where necessary for improved treatment of neuropsychiatric disorders. AAU15114 necessary for improved treatment of neuropsychiatric disorders. AAU15114 is soforms used in the method of the invention
Diagnosing and monitoring Schizophrenia by detecting the presence of Schizophrenia Associated Features and Schizophrenia Associated Protein Isoforms in samples of cerebrospinal fluid.
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28-DEC-2000; 2000US-00750395.
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밁 Ś Matches Query Match Best Local & 9 2 FGYKN 6 4: Similarity FGYSN 13 Conservative 49.0%; Score 25; DB Pred. No. 9.8e 0 ;:)B :, 9.8e+02; 1; DB 4; Length 14; Indels 0; Gaps 0

Sequence 14 AA;

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RESULT 39
ABB52298
Human; neuroprotective; nootropic; gene therapy; vaccine; Alzheimer's disease; Alzheimer's Disease-Associated Feature; Alzheimer's Disease-Associated Protein Isoform; API; tryptic
                                                                                                                                            Human API-84
                                                                                                                                                               08-FEB-2002
                                                                       Homo sapiens
                                                                                                                                                                                                      ABB52298 standard; peptide; 14
                                                                                          Expression Reference Protein Isoform; ERPI;
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                                                   WO200175454-A2
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digest;
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03-APR-2001; 2001WO-US010908

11-OCT-2001.

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RESULT 40
AAE25488
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AC AAE25
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AC CAPI-
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Chron
KW Chron
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PD 13-JU
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PD 10-DE
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PT Garee
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to methods for the screening, diagnosis and prognosis of Alzheimer's disease. The methods involve the detection of Alzheimer's Disease-Associated Features (AFs) and Alzheimer's Disease-Associated Protein Isoforms (APIs) in cerebrospinal fluid, serum or plasma. The abundance of the AFs and APIs is then normalised to an Expression Reference Protein Isoform (ERPI) in order to determine whet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Screening for Alzheimer's disease in a mammal, by making two-dimensional array of a feature whose relative abundance correlates with disease, and comparing with abundance of the feature in samples of healthy persons.
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Potter
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                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
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28-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          asthma-associated asthma-associated peptide; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tryptic
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PFIZER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        It is suffering from, or has a predisposition to Alzheimer's The relative abundance of the AFs and APIs correlates with the of Alzheimer's Disease. The present sequence is a peptide from an API by proteolysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14
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Rohlff C,
Rohite F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA;
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                                                                                                                                                                                     GLYCOSCIENCES
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                                                                                                                                        JA;
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80.
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, Silber BM, Sti
F, Williams SA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein isoform; CAPI; CAF; chronic asthma;
feature; screening; diagnosis; prognosis;
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9.8e+02;
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Screening chronic asthma by analyzing a test tissue sample by to dimensional electrophoresis to generate a two dimensional array features and comparing the abundance of the test sample with a second comparing the second comparin

a sample t wo 2002-537580/57

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RESULT 41
ABJ15069
ID ABJ1
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AC ABJ1
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AC ABJ1
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Best Local S
Matches 4
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                 The present invention relates to compositions comprising a substance that modulates the status of 125P5C8 or a molecule that is modulated by 125P5C8. The status of a cell that expresses 125P5C8 is modulated. The composition is useful for treating cancer, particularly prostate, bladder, kidney, colon, ovary or breast cancer. The 125P5C8 protein and/or a nucleotide sequence encoding the protein is useful for immunising a mammal against cancer. The present sequence is a 125P5C8 epitope shown in the exemplification of the invention
                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                           New composition comprising a substance that modulates the status 125P5C8 gene or a molecule that is modulated by 125P5C8, useful 1 treating or preventing cancer that expresses or over expresses 12
                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
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RK, Morrison
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kidney cancer; colon cancer; ovari
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No. 9.8e+02;
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                                                                                                                                                                                                                                                                                                               he status of
, useful for
presses 125P5C8.
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                                                                            cc screening, diagnosis or prognosis of Alzheimer's disease (AD) in a cc subject. The method comprises analysing a sample of brain tissue from a cc subject by 2D electrophoresis to generate a 2D array of Alzheimer's cc disease-associated features (ADFs), whose relative abundance correlates cc with the presence, absence, stage or severity of AD and comparing the cc abundance of each feature with the abundance of that chosen feature in cc brain tissue from persons free from AD. The invention also describes cc Alzheimer's disease-associated protein isoforms (ADPIs) detectable in cc brain tissue. The methods and compositions of the invention are useful cc determining the stage or severity of AD in a subject, for cd determining the stage or severity of AD in a subject, for identifying a composition are useful for treating or preventing AD, and for determining to a subject having AD. Antibodies capable of binding to efficacy of a given treatment regime. An agent that modulates the cc activity of ADPI is useful in the manufacture of a medicament for the treatment or prevention of AD in a subject. ABG67292-ABG68038 represent comman ADPI tryptic digest peptides
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                                                                                                                                                                                                                                                                                                                                                                                  Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; Alzheimer's disease; AD; brain tissue; ADF; ADPI;
Alzheimer's disease-associated feature; neuroprotective;
Alzheimer's disease-associated protein isoform; nootropi
                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (OXFO-) OXFORD GLYCOSCIENCES
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Pred. No. 9.8e+02;
2; Mismatches 3
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               DB 5;
9.8e+02
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7.8e+02;
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ABR59105 standard; peptide;

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                                                                                                                                     The present invention describes an isolated salivary polypeptide (I), of phlebotomus papatasi. Also described: (1) an isolated nucleic acid (II) encoding (I); (2) an isolated fragment (III) of (II), where the fragment CC encodes the polypeptide fragment specific for a polypeptide such as CC phlebotomus papatasi salivary polypeptide (PpSP) 12, PpSP14, PpSP15, CC ppSP30, PpSP36, PpSP42 or PpSP44; (3) an isolated antigenic or immunogenic fragment (IV) of (I); (4) a nucleic acid (V) that hybridises (III); and (6) a composition (VII) comprising (I) or its fragment, (IV) or (III); and (6) a composition (VII) comprising (I) or its fragment, (IV) and immunostimulant activities, and can be used in vaccines. (VII) is useful for producing an immune response in a subject or for preventing Leishmaniasis in a subject. (II) is also useful for preventing Leishmaniasis in a subject. (VII) is useful as a vaccine. ACC79987 to ACC79999 and ABR91176 to ABR92046 represent sequences used in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated salivary polypeptide of Phlebotomus papatasi, producing an immune response in a subject or for preventing
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Synthetic.
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protozoacide;
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                                                                                                               Sequence 14 AA;
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                                                                     49.0%;
                                                         Score 25; DB Pred. No. 9.8e
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vaccine; immune response; Leishmaniasis.
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Leishmaniasis
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Potter DM,
Sunderland
                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to methods for screening or diagnosing Alzheimer's disease (AD) to determine the stage or severity of AD in a subject, to identify subject at risk of developing AD, or to monitor the effect of therapy administered. The methods comprise analysing a test sample of body fluid by 2-dimensional electrophoresis to generate a 2-dimensional array of AD-associated features (AFs). The method alternatively comprises quantitatively detecting in a sample of body fluid from the subject, one or more AD-associated protein isoforms (APIs, approximate the subject, one or more AD-associated protein isoforms (APIs)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Screening or diagnosing of Alzheimer's disease (AD) determine the stage or severity of AD in a subject, comprises analyzing a test sample of body fluid from the subject by 2-dimensional electrophoresis.
                                                       schizophrenia; Parkinson's dise
Alzheimer's disease-associated
                                                                                                                 Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 14
 US2003064411-A1
                            Homo sapiens
                                                                    human; Alzheimer's disease; vascular dementia; Lewy body schizophrenia; Parkinson's disease; multiple sclerosis; d
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Rohlff C, Silber BM, Snyder
PT, Townsend RR, White WF, W
                                                                                                             disease-associated protein isoform tryptic peptide #462
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er PJ, Soares HD,
Williams SA;
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                                                                                    dementia;
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                                                                                                                                               The method is also useful in treating vascular dementia, Lewy body dementia, schizophrenia, Parkinson's disease, multiple sclerosis or depression. The inventive method identifies sensitive and specific biomarkers for the diagnosis of Alzheimer's disease in living subjects. It provides therapeutic agents for Alzheimer's disease that works quickly, potently, specifically with fewer side effects. The present sequence represents the amino acid sequence of a Alzheimer's disease-associated protein isoform tryptic peptide.
                                                                                                                                                                                                                                                                          The invention relates to a method of screening or diagnosing Alzheimer's disease in a subject. The method is useful for screening, diagnosis or prognosis of Alzheimer's disease in a subject for determining the stage of severity of Alzheimer's disease in a subject, for identifying a subject at risk of developing Alzheimer's disease, or for monitoring the effect of therapy administered to a subject having Alzheimer's disease.
                                                                                                                       Sequence 14 AA;
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(PARE/)
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Q8hu93 bryum donia
Q50268 phytoplasma
Q9tu33 canis famil
078121 oreochromis
Q9twm3 locusta mig
Q865c9 sus scrofa
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Q53579 rhodobacter
Q8n6b2 homo sapien
P87031 saccharomyc
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Q9TQV4;
Q9TQV4;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
POP-variant * transferrin gene, EXON 17 and partial
                                                                                                                                                                                                                                                                                    EMBL; AI
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Giffard J.M., Brandon R.B., Bell T.K.;

"Further identification of single nucleotide polymorphisms
equine transferrin gene.";
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Perissodactyla; Equidae; Equus.
                                                                                                                                                                                                                                                  SEQUENCE
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FHRV 10
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Pred. No.
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01-JUL-1997
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Q53579;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Capacitative calcium channel protein Trpl (Fragment).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-OCT-2002 (TrEMBLrel. 22, 01-OCT-2002 (TrEMBLrel. 22, 01-OCT-2002 (TrEMBLrel. 22,
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MEDLINE=92234963; PubMed=1569029;
Richter P., Brand M., Drews G.;
"Characterization of LHI- and LHI+
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rhodobacter capsulatus (Rhodopseudomonas capsulata).
Bacteria; Proteobacteria; Alphaproteobacteria; Rhodobacterales;
Rhodobacteraceae; Rhodobacter.
                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
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MEDLINE=22181008; PubMed=12193412;
Yang M., Gupta A., Shlykov S.G., C
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NCBI_TaxID=1061;
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SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

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STRAIN=BALB/c; TISSUE=Kidney;

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STRAIN=BALB/c; TISSUE=Kidney;

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STRAIN=BALB/c; TISSUE=Kidney;

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ORF YGR126W (Fragment).
Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomyces.
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                  Submitted (JUL-2002) to the EMBL; AF530457; AAN28710.1; NON_TER 1 1 1 NON_TER 12 12 SEQUENCE 12 AA; 1448 MW;
                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=BALB/c; TISSUE=Kidney;
MEDLINE=20334280; PubMed=10873615;
Alliel P.M., Seddiqi N., Goudou D.,
Velasco E., Rieger F., Perin J.P.;
"Myoneurin, a novel member of the B1
expressed in human muscle.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Van Dyck L., Skala J., de We
Nawrocki A., Del Bino S., Go
Submitted (MAY-1996) to the
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Eukaryota; Metazoa; (
Mammalia; Eutheria; )
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3; Conser
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 KNLKKHK
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                                                                          Conservative
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Rodentia;
                                                                                             35.0%;
57.1%;
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75.0%;
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S., Goffeau A.;
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Pred.
1; Mis
                                                                           Score 21; DB Pred. No. 3.9e 1; Mismatches
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Last sequence update)
Last annotation update)
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AC Q8
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Best Local S
Matches 4
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Best Local S
Matches 2
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Q9UNM9;
01-MAY-2000
Q8HUB4;
Q8HUB4;
01-MAR-2003
                                                                                                                                                                                                                                                                                                  056127
056127;
                                                                                                                    SEQUENCE FROM N.A.
STRAIN=pmws PCV;
Hamel A.L., Lin L.L., Nayar G.P.S.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ
EMBL; AF027217; AAC59472.1; -.
SEQUENCE 14 AA; 1761 MW; 00F50DC89EBAF20B
                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AF107699; AAD24963.1; -...
InterPro; IPR001487; Bromodomain.
NON_TER 1 1 1
NON_TER 14 14
                                                                                                                                                                           SEQUENCE FROM N.A.
STRAIN=pmws PCV;
STRAIN==98241772; PubMed=9573301;
Hamel A.L., Lin L.L., Nayar G.P.;
"Nucleotide sequence of porcine circovirus
multisystemic wasting syndrome in pigs.";
J. Virol. 72:5262-5267(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                    Porcine circovirus.
Viruses; ssDNA viruses;
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                                                                                                                                                                                                                                           NCBI_TaxID=46221;
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Keddache M, Zhang F, Greenberg
"A Dinucleotide Repeat Between
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                                                                                     Similarity 4; Conserv
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(Tremblie).
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(TrEMBLrel.
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Primates;
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23,
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Last sequence
Last annotation
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Last
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Pred. No. 6.9e
1; Mismatches
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Pred. No. 6.9e
3; Mismatches
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Exons 2
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annotation update)
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annotation
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6.9e+03;
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RESULT 9
Q8HU93
Q8HU93
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DT 01-M
DT 01-M
DT 01-J
DE Ribo
GN RPL1
OS Bryu
OG Chlo
OC Bryo
OC NCBI
RN [1]
RP SEQU
RC STRA
RA Pede
RL SUBT
DR COBJ
DR GO;
KW Chlc
FT NON
FT NON
SEQU
RESULT 10
Q50268
ID Q5026
AC Q5026
DT 01-NO
DT 01-JA
DT 01-DE
Rp116
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Best Local S
Matches 3
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Best Local S
Matches 3
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Pedersen N., Cox C., Hedenas L.;

Submitted (SEP-2002) to the EMBL
EMBL; AF546765; AAN62980.1; -.

GO; GO:0009507; C:chloroplast; T:

Chloroplast.

NON_TER
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NO
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Eukaryota; Viridiplar
Bryopsida; Bryidae; E
NCBI TaxID=67232;
                      Q50268;
Q50268;
01+NOV-1996
01-JAN-1999
01-DEC-2001
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01-MAR-2003
01-JUN-2003
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NON_TER
SEQUENCE
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Eukaryota; Viridiplantae; Streptophyta; Embry Bryopsida; Bryidae; Bryales; Bryaceae; Bryum. NCBI_TaxID=66993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=BD628;
Pedersen N., Cox C., Hedenas L.;
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ
EMBL; AF546786; AAN63001.1; -.
GO; GO:0009507; C:chloroplast; IEA.
Chloroplast.
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Bryidae; Bryales;
                     (TrEMBLrel. 01, 09) (TrEMBLrel. 09, 00) (TrEMBLrel. 19, 00)
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42.9%;
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ss; Bryaceae; Anomobryum.
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EMBL/GenBank/DDBJ
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7.6e+03;
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9.1e+03;
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RESULT 12
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AC 07812
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DT 01-DE
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O78121;
O1-NOV-1998
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Phytoplasma sp.

Bacteria; Firmicutes; M
Acholeplasmataceae; Phy
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MEDLINE=20154965; PubMed=10690375;
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Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; C
Mammalia; Eutheria; Carnivora;
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Q9TU33;
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EMBL; M74770; AAA25331.1; -.

NON_TER 14 14

SEQUENCE 14 AA; 1870 MW; 87C31:
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"DNA sequence of the ribosomal protein genes plant-pathogenic mycoplasma-like organism.";

""" wirrobiol. Lett. 84:71-74(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQÜENCE
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Lim P.-O., Sears B.
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Genet. 31:76-77(2000).
AF159258; AAD56289.1;
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; Fissipedia; Canidae;
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Glutamine synthetase (
Sus scrofa (Pig).
Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Ce
NCBI_TaxID=9823;
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01-JUN-2003
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MEDLINE=98315113; PubMed=9649539;

Malaga-Trillo E., Zaleska-Rutczynska Z., McAndrew B., Vincek V.,

Malaga-Trillo E., Zaleska-Rutczynska Z., McAndrew B., Vincek V.,

Figueroa F., Sultmann H., Klein J.;

Figueroa F., Sultmann H., Figueroa F., Sultmann H., Figueroa F., Sultmann H., Figueroa F., Sultma
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SEQUENCE
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MEDLINE=95063174; PubMed=7972937;
Schoofs L., Veelaert D., Holman G.M.,
"Partial identification, synthesis and
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Oreochromis niloticus (Nile tilapia) (Tilapia nilotica)
Dreochromis niloticus (Nile tilapia) (Tilapia nilotica)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Ne
Acanthomorpha; Acanthopterygii; Percomorpha; Perciforme
Cichlidae; Oreochromis.
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridoidea; Acrididae; Oedipodinae; Locusta.
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01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
LOCUSTAMYOINHIBIN, LOM-MIH-MYOINHIBITING neuropeptide.
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Locusta migratoria.";
Regul. Pept. 52:139-156(1994).
SEQUENCE 13 AA; 1464 MW; D
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                          Chordata; Craniata; Vertebrata;
Cetartiodactyla; Suina; Suidae;
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1.5e+04;
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Euteleostei; Neoteleostei;
rpha; Perciformes; Labroidei;
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Submitted (NOV-2000) to Swiss-Prot.
-!- FUNCTION: BINDS TO THE VCAM-1 5'UTR
-!- SIMILARITY: TO THE FETUIN FAMILY.
PIR; A59387; A59387.
GO; GO:0003723; F:RNA binding; IEA.
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Kim J.G., Vallet J.L., Christenson R.K.;

"Characterization of porcine glutamine synthet

Submitted (JAN-2003) to the EMBL/GenBank/DDBJ.

EMBL; AY216477; AA064254.1; -.
                                                                                       Pisum sativum (Garden pea).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids;
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PS01255; FETUIN_2;
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A Peltier J.-B., Friso G., Kalume D.E., Roepstorff P., Nilsson A Adamska I., van Wijk K.J.;
A Adamska I., van Wijk K.J.;
"Proteomics of the chloroplast: systematic identification an repretent analysis of lumenal and peripheral thylakoid protectargeting analysis of lumenal and peripheral thylakoid; IEA.

GO; GO:0009579; C:chloroplast; IEA.

GO; GO:0009579; C:chloroplast; IEA.

GO; GO:0009579; C:chloroplast; IEA.
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Q8CG13;
01-MAR-2003
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01-MAY-2000
01-JUN-2002
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"Binding site mapping of a photoaffinity-labeled binding protein.";
Biochem. Biophys. Res. Commun. 182:466-473 (1992)
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Glutamate receptor ionot
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia; Sph
Sphingidae; Sphinginae; Manduca.
NCBI_TaxID=7130;
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Mus musculus (Mouse
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rmone binding protein, JHBP=4.9 kDa GI
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MEDLINE=98027386; PubMed=9361435;
Medline=7 Tiebert C.A., Smith T.,
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01-NOV-1999
01-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=97208220; PubMed=9055422; Liebert C.A., Wireman J., Smith T., Summers A.O.; "Phylogeny of mercury resistance (mer) operons of gram-negative bacteria isolated from the fecal flora of primates."; Appl. Environ, Microbiol. 63:1066-1076(1997).
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                                                                                                                                  SEQÜENCE
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[3]
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Enterobacteriaceae; Escher
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NCBI_TaxID=10090;
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          "Genetic mapping of the lipid transfer radiata and Pinus taeda."; Submitted (MAR-2001) to the EMBL/GenBan EMBL; AY029028; AAK40266.1; -. NON_TER 11 11
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                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
STRAIN=PRLTP9-ii58055;
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MEDLINE=97208220; PubMed=9055422;
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Bacteria; Proteobacteria;
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Jiebert C.A., Watson A.L.. Summers
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AY029028; AAK40266.1; -.
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Q9TT31;
01-MAY-2000
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SEQUENCE
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Q9TT29;
O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Callitrichidae; Saguin
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Mammalia; Eutheria; Primates; Platyrrhini; Callitrichidae; Callit
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01-MAY-2000 (TrEMBLrel. 13,
01-DEC-2001 (TrEMBLrel. 19,
Beta-globin (Fragment).
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Mol. Biol. Evol. 17:416-422(2000).
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Francino M.P., Ochman
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Pred. No. 1.9e
1; Mismatches
                                                      Score 17; DB
Pred. No. 2.1e
2; Mismatches
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Pred. No. 2.1e+04;
2; Mismatches '
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                                                                                         DB 6;
2.1e+04;
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1.9e+04;
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RESULT
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Q7ZZN6
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01-JUN-2003
01-JUN-2003
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Zinc finger E
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01-NOV-1996
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Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
Enterobacteriaceae; Escherichia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; MON_TER
                                                                                                         "Identification of a phylogenetically conserved activin-responsive enhancer in the Zic3 gene.";
Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AF506278; AAP20809.1; -.
                                                                                                                                                                                                   Eukaryota; Metazoa; C
Amphibia; Batrachia;
Xenopodinae; Xenopus.
                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                          "A family of regulatory genes associated modification systems.";
J. Bacteriol. 173:1367-1375(1991).
EMBL; M63620; AAA24557.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. MEDLINE=91139577; I
                                                                               SEQUENCE
                                                                                                                                                                                                              Xenopus laevis (African clawed frog).
Eukaryota; Metazoa; Chordata; Craniata; Vej
Amphibia; Batrachia; Anura; Mesobatrachia;
                                                                                                                                                                                                                                                                                                        Q7ZZN6
                                                                                                                                                   Weber J
                                                                                                                                                            SEQUENCE FROM N.A. TISSUE=Cerebellum;
                                                                                                                                                                                          NCBI_TaxID=8355;
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KHMKVH
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(TrEMBLrel. 24, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
protein (Fragment).
                                       Conservative
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C., Blumenthal F
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                                               28.3%;
50.0%;
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                                     Score 17;
Pred. No.
1; Mismatc
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Pred. No.
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                                                 2.3e+04;
                                                          DB 13;
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ia; Pipoidea; Pipidae;
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Porcine circovirus type 2-
Porcine circovirus type 2-
Porcine circovirus type 2-
Porcine circovirus type 2-
Porcine circovirus, and
Porcine circovirus type 2.
Viruses; ssDNA viruses; Ci
NCBI_TaxID=46221, 86385, 8
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O93202; Q9YJ34;
O1-NOV-1998 (TrEMBLrel. 0
01-NOV-1998 (TrEMBLrel. 0
01-OCT-2003 (TrEMBLrel. 2)
                                                                                            SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

SPECIES=porcine circovirus type 2; STRAIN=24657 NL;

SPECIES=porcine circovirus type 2; STRAIN=24657 NL;

SPECIES=porcine circovirus P.J.G.M.,

Wellenberg G.J., Pesch S., Berndsen F.W., Steverink P.J.G.M.,

Hunneman W., 'Van der Vorst T.J.K., Peperkamp N.H.M.T., Ohling,

Schippers R., Van Oirschot J.T., de Jong M.F.;

Schippers R., Van Oirschot J.T., de Jong M.F.;

"Isolation and characterization of porcine circovirus type 2

"howing signs of post-weaning multisystemic wasting syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         [3]
SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
SPECIES=porcine circovirus type 2-D;
SPECIES=porcine circoviruses.";
Hamel A.L., Nayar G.P.S.;
Hamel A.L., Nayar G.P.S.;
"Genetic characterization of four novel type-2 Porcine circoviruses.";
                                                      EMBL;
                                                                                                                                                                                                                               SPECIES=bovine circovirus;
Hamel A.L., Nayar G.P.S.;
"Nucleotide sequence of a circovirus detected clinical syndromes.";
   EMBL;
                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.

SPECIES-porcine circovirus type 2-B;

Hamel A.L., Nayar G.P.S.;

"Nucleotide sequence of four different isolates of porcine detected in pigs with various clinical syndromes.";

Submitted (DEC-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SPECIES=porcine circovirus; STRAIN=PORCINE CIRCOVIRUS MEDLINE=98418498; PubMed=9747726; Meehan B.M., McNeilly F.M., Todd D., Kennedy S., Jewhu Ellis J.A., Hassard L.E., Clark E.G., Haines D.M., All "Characterization of novel circovirus DNAs associated syndromes in pigs."; J. Gen. Virol. 79:2171-2199(1998).
                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES=porcine circovirus type 2-C, and Hamel A.L., Nayar G.P.S.;
"Nucleotide sequence of four different in pigs with various clinical syndromes. Submitted (NOV-1998) to the EMBL/GenBank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SPECIES=porcine
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 AF055394;
AF055391;
AF055392;
AF055393;
AF117753;
AF117753;
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 AAC35336.1;
AAC35305.1;
AAC35316.1;
AAC35326.1;
AAD12313.1;
AAD12313.1;
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95543, 85544,
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Haines D.M., Allan O
DNAs associated with
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, 85709, 85542,
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r., Ohlinger V.
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with wasting
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RESULT
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DT 011
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Best Local
SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN=PWB;
STRAIN=PWB;
Lee I.-M., Martini M., Marcone C., Zhu S.;
Lee I.-M., Martini M., Marcone C., Zhu S.;
Lee I.-M., Martini M., Marcone C., Zhu S.;

"Classification of phytoplasma strains in the elm yellows group

"Classification of phytoplasma ulmi for the

(16SrV) and proposition of 'Candidatus Phytoplasma

"he phytoplasma associated with cherry lethal yellowing and the cherry lethal yellowing and 
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q7X4C1
Q7X4C1;
01-0CT-2003
01-0CT-2003
01-0CT-2003
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01-NOV-1996
01-NOV-1996
01-JUN-2003
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NON TER
SEQUENCE
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EMBL; AF112862; AAD03090.1;
EMBL; AF109397; AAD11937.1;
EMBL; AF201897; AAG41230.1;
SEQUENCE 14 AA; 1747 MW;
                                                                                                                                                                                                                                                                                                             Potato witches'-broom phytoplasma.
Bacteria; Firmicutes; Mollicutes;
Acholeplasmataceae; Phytoplasma.
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                                                                                                                                                                                                                                                                              Acholeplasmataceae;
NCBI_TaxID=37701;
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MEDLINE=88015611; PubMed=3658701;
Portner S.G., Yanofsky M.F., Nest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=356;
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"Isolation of a Precisely Delineated,

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Submitted (JUN-2002) to the EMBL; AY123324; AAM96898.1;
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Cotter J.-P., StJohn J., Quinn T.W.;
"Isolation of a Precisely Delineated,
Repeat 1 (CR1) Retrotransposon Within
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Coscoroba.
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                                         CRC64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ebrata; Euteleostomi;
Anatidae; Anser.
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n of the Coscoroba
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                                                                                                                                                                                                                         RESULT 32
Q8AWV7
AC Q8AWV
AC Q8AWV
DT 01-MA
DT 01-MA
DT 01-MA
DT 01-MA
DT 01-MA
COLUBRAT
OC ARCHOO
OX NCBI
RN [1]
RP SEQUE
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Q8AWV8
AC Q8AWV8
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Best Local Similarity
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8 HRVI 11
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Query Match
Best Local S
Matches 1
                                                                                                                         SEQUENCE FROM N.A.

Cotter J.-P., StJohn J., Quinn T.W.;

Cotter J.-P., StJohn J., Quinn T.W.;

"Isolation of a Precisely Delineated,

"Isolation fa Precisely Delineated,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; AY1
NON_TER
NON_TER
SEQUENCE
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01-MAR-2003
01-MAR-2003
01-MAR-2003
                                                                Repeat 1 (CR1) Retrotransposon coscoroba.";
Submitted (JUN-2002) to the EME EMBL; AY123327; AAM96901.1; -.
NON TER 1 1
NON_TER 8 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (JUN-2002) to the Submitted (AMP) to the Submitted (AMP) 1
                                                                                                                                                                                                                                                                        01-MAR-2003 (TrEMBLrel.
01-MAR-2003 (TrEMBLrel.
01-MAR-2003 (TrEMBLrel.
                                                                                                                                                                                                                       Anas platyrhynchos (Domestic duck).
Eukaryota; Metazoa; Chordata; Crani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cygnus columbianus (tundra swan).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euk
Archosauria; Aves; Neognathae; Anseriformes; Anatidae;
NCBI TaxID=110926;
                                                                                                                                                                                           NCBI_TaxID=8839;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.

Cotter J.-P., StJohn J., Quinn T.W.;

"Isolation of a Precisely Delineated,

Repeat 1 (CR1) Retrotransposon Within
                                                                                                                                                                                                                                                               LDH-B (Fragment).
                                                                                                                                                                                                          Archosauria; Aves; Neognathae; Anseriformes;
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LDH-B.
l Similarity
1; Conserv
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1; Conserv
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nilarity 25.0%;
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  Conservative
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              26.7%;
25.0%;
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                                                   MW;
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Last sequence update)
Last annotation update)
Score 16; DB
Pred. No. 1e+0
3; Mismatches
                                                                                                      EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                        Created)
Last sequence update)
Last annotation updat
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Pred.
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No. 1e+06;
           DB 13;
1e+06;
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Q9UR95
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Q8MEL7
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Q8MEM2
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Best Local S
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Best Local S
Matches 2
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Q9UR95;
01-MAY-2000
01-MAY-2000
01-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GO; GO: 000750
Chloroplast.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-2002
01-OCT-2002
01-JUN-2003
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NON_TER
SEQUENCE
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Evers M.E., Huhse B., Titorenko V.I., Kunau W.H
Harder W., Veenhuis M.;
"Affinity purification of molecular chaperones
polymorpha using immobilized denatured alcohol
FEBS Lett. 321:32-36(1993).
SEQUENCE 11 AA; 1230 MW; 71872C1779C3372B C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chloroplast.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Malvales; Malvaceae; Malvoideae; Sida.
Q8MEM2;
01-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chloroplast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. pfeil B.E., Brubaker C.L., "phylogeny of Hibiscus and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sida hookeriana.
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01-MAY-2000 (TrEMBLrel. 13, La
01-JUN-2000 (TrEMBLrel. 14, La
Heat shock protein 60 homolog
                                                            Q8MEM2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oroplast DNA sequences of not. 27:333-350(2002).
L; AF384624; AAM50396.1; -.
GO:0009507; C:chloroplast;
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2; Conserv
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3; Conserv
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11 AA;
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(TrEMBLrel. 22, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
cotein 16 (Fragment).
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                                                                                                                                                                                                                                                                                                                                       Conservative
     (TrEMBLrel.
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                                                                 PRELIMINARY,
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50.0%;
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28.6%;
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the tribe Hibisceae
of ndhF and the rpl16
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Pred. No.
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Pred. No.
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l; Mismatches
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3e+04;
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bisceae (Malvaceae)
the rpl16 intron.";
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RESULT 38
Q8MES5
ID Q8MES
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DT 01-OC
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DE Ribos
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OG Chlor
OC Eukar
OC Eukar
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Chlo:
GN GO;
KW Chlo:
RT Syst
DR GO;
KW Chlo:
RT Syst
DR GO;
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RESULT 39
Q8MEP0
ID Q8MEP
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EMBL;
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01-JUN-2003
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01-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pfeil B.E., Brubaker C.L., Craven L.A., Crisp M.D.; "Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) chloroplast DNA sequences of ndhF and the rpl16 intron.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. Pfeil B.E., Brubak
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Chloroplast
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L; AF384561; AAM50399.1; -.
GO:0009507; C:chloroplast;
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nilarity 28.6%;
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annotation update)
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(Malvaceae) using
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Pfeil B.E., Brubaker C.L., Craven L.

Pfeil B.E., Brubaker C.L., Craven L.

Phylogeny of Hibiscus and the tribe chloroplast DNA sequences of ndhF ar Syst. Bot. 27:333-350(2002).

EMBL; AF384566; AAM50404.1; -.

GO; GO:0009507; C:chloroplast; IEA.

Chloroplast.
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01-OCT-2002
01-JUN-2003
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NCBI_TaxID=183226;
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EMBL; AF384598; AAM50370.1; -...GO; GO:0009507; C:chloroplast; IEA.
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el. 22, Last sequence up
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(Fragment).
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                                                                                                                                                                                        ; Score 16; DB; Pred. No. 3e+0
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RESULT 43
Q8MEL9
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OC Chlor
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01-OCT-2002 (TrEMBLrel. 2:
01-OCT-2002 (TrEMBLrel. 2:
01-JUN-2003 (TrEMBLrel. 2:
                                                                                                                                                                                                                          Pfeil B.E., Brubaker C.L., Crave "Phylogeny of Hibiscus and the t chloroplast DNA sequences of ndh Syst. Bot. 27:333-350(2002).
EMBL; AF384581; AAM50353.1; -. GO; GO:0009507; C:chloroplast; I
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NCBI TaxID=183253;
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eurosids II; Malvales;
NCBI_TaxID=183239;
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Hibiscus normanii.
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01-OCT-2002 (TrEMBLrel. 22, Last
01-JUN-2003 (TrEMBLrel. 24, Last
Ribosomal protein 16 (Fragment).
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the tribe Hibisceae (Malvaceae) using
of ndhF and the rpl16 intron.";
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EMBL; AF384622; AAM50394.1; -. GO; GO:009507; C:chloroplast; I
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid
eurosids II; Malvales; Malvaceae; Malvoideae; Hibiscus.
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Phylogeny of Hibiscus and
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t. Bot. 27:333-350(2002).
L; AF384578; AAM50416.1; -.
GO:0009507; C:chloroplast;
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the tribe Hibisceae (Malvaceae)
of ndhF and the rpl16 intron.";
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the tribe Hibisceae
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DT 01-CUT-2002 (TrEMBLrel. 22, Last sequence update)

DE RIASYOTA; PROMINER MSD 11 AA.

CC PROPERTY MSD 11 AA.

PRELIMINER PROMINER C.L., Craven L.A., Crisp M.D.;

RT "Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) using RT Chloroplast DA.

RA Pfeil B.E., Brubaker C.L., Craven L.A., Crisp M.D.;

RT "Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) using RT Chloroplast DA.

RA Pfeil B.E., Brubaker C.L., Craven L.A., Crisp M.D.;

RT "Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) using RT Chloroplast DA.

RA Pfeil B.E., Brubaker C.L., Craven L.A., Crisp M.D.;

RT "Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) using RT Chloroplast DA.

RA Pfeil B.E., Brubaker C.L., Craven L.A., Crisp M.D.;

RT "Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) using RT Chloroplast DA.

RA Pfeil B.E., Brubaker C.L., Craven L.A., Crisp M.D.;

RT "Phylogeny of Hibiscus and the tribe Hibisceae (Malvaceae) using RT Chloroplast J. T.

RE EMBL, AR304563; AAM50401.1; -.

DR CO, GO:,0005507; C:chloroplast; IEA.

WON TER 11 AA; 1470 MM; 7227C35ID32409D4 CRC64;

CO GO:,0005507; C:chloroplast; IEA.

WON TER 11 AA; 1470 MM; 7227C35ID32409D4 CRC64;

Matches 2; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

DB SEQUENCE C.L., Craven L.A., Crisp M.D., Cri
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Copyright (c) 1993 - 2004 Compugen Ltd.
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NP1_LYMST
NP4_LYMST
TEML_RANTE
RS19_PPWBP
SMS1_MYOSC
SMS_ALLMI
FAR4_CALVO
                                                 FAR7_MACRS
MY14_EISFO
LCK6_LEUMA
ASL1_BACSE
CXL1_CONMR
CXL4_CONMR
FARP_CALSI
COXM_RAT
LCX4_PHEVI
LCK4_LEUMA
FAR2_PEUMO
TKL4_LOCMI
ANGT_HORSE
RS19_CLOPP
RS1_ERWCH
ULAD_HUMAN
TKN1_SCYCA
CA42_LITCI
TKC2_CALVO
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FAR7_PENMO
LMA1_LOCMI
COXG_RAT
                                                                                                                                                                                                                                                               SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                       585
                                                                                                                                                      P80430 rattus norv
P83280 macrobrachi
P46979 eisenia foe
P19988 leucophaea
P83146 bacteroides
P58807 conus marmo
P58810 conus marmo
P38495 callinectes
 P30250
P01016
Q446228
Q446228
P37985
P37985
P31929
P031929
P031929
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P60118
P80118
P80118
P87108
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P87108
P31885
P31885
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P21143
P83317
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P83322 penaeus mon
P38496 locusta mig
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      8 lymnaea sta
1 lymnaea sta
4 rana tempor
3 pigeon pea
0 myoxocephal
                                                 9 homo sapien
8 scyliorhinu
2 litoria cit
8 calliphora
                                                                           B clover prol
B loofah witc
S erwinia chr
                                                                                                7 penaeus mon
0 locusta mig
5 equus cabal
                                                                                                                    pseudonaja
pheretima v
leucophaea
                                                                                                                                          rattus norv
RESULT 2
FAR7_PENMO
ID FAR7_PENMO
AC P83322;
DT 28-FEB-2003
DT 28-FEB-2003
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| al Similarity 42.9%; Preside Similarity 42.9%; President S | BCELLULAR LOCATION: Secre MILARITY: Belongs to the 61033; B61033. ro; IPR002040; Tachy Neur E; PS00267; TACHYKININ; 1 inin; Neuropeptide; Amida S 10 10 AA; 1210 MW; 91 CE 10 AA; 1210 MW; 91 | PubMed=8210506 H., Hino J., inins in frog -88(1993). ykinins are ac al responses, and contract | CE, AND SYNTHESIS. =Brain; E=91254337; PubMed=204314 H., Hino J., Minamino N. tion of four novel tachyk and intestine."; m. Biophys. Res. Commun. | (Rel. 19, Create (Rel. 19, Last s (Rel. 42, Last a iin B (RTK B). siana (Bull frog) fetazoa; Chordata (trachia; Anura; 1410; | STANDARD; P | 14 23.3 9 1 FIBB 14 23.3 9 1 HUTU 14 23.3 10 1 ANGT 14 23.3 10 1 PVK I 14 23.3 10 1 TRP8 14 23.3 10 1 TKN I 14 23.3 11 1 TKN I 14 23.3 12 1 LICH 14 23.3 13 1 DHC 14 23.3 13 1 NEUT 14 23.3 13 1 NEUT 14 23.3 13 1 NEUT 14 23.3 14 1 ADFA |
| red. No. 3e+02; Mismatches 2; Indel | hykinin famil nin. nin. TION. 56B59D5BAB5 C | Minamino N., Matsuo H.; Minamino N., Matsuo H.; (Rana catesbeiana) brain ctive peptides which excit are potent vasodilators a (directly or indirectly) | ; Kangawa K., Matsuo H nins from frog (Rana 77:588-595(1991). | update) on update) ata; Vertebrata; achia; Ranoidea; | ALIGNMENTS | FIBB PAPHA HUTU KLEAE ANGT BOVIN PVK LOCMI RL16 ACHLA TRP8 LEUMA TKN PHYFU LICH BACLI XYLA STRVN IDHC PIG NEUT CAVPO ADFA TENMO |
| els 0; Gaps 0; | • | ain and ain and xcite neurons, rs and ly) many smooth | catesbeiana) | Euteleostomi; Ranidae; Rana. | | P19343 papio hamad P12381 klebsiella P01017 bos taurus P01017 bos taurus P83382 locusta mig P29221 acholeplasm P81740 leucophaea P08615 physalaemus P82907 bacillus li P82907 bacillus li P14405 streptomyce P20304 sus scrofa P32560 cavia porce P82965 tenebrio mo |

(Rel. 41, Created)
(Rel. 41, Last sequence update)

STANDARD;

PRT;

12 AA.

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RESULT 3
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P38496;
01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
Lom-AG-myotropin I (Accessory gland myotropin I).
Locusta migratoria (Migratory locust).
Locusta; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Metazoa; Orthoptera; Caelifera; Acridomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE 21956277; PubMed=11959015;
Sithigorngul P., Pupuem J., Krungkasem C., Longyant Chaivisuthangkura P., Sithigorngul W., Petsom A.;
"Seven novel FMRFamide-like neuropeptide sequences of the giant tiger prawn Penaeus monodon.";
Comp. Biochem. Physiol. 131B:325-337(2002).
-!- SUBCELLULAR LOCATION: Secreted.
-!- MASS SPECTROMETRY: MW=1381.4; METHOD=MALDI.
-!- SIMILARITY: Belongs to the FARP (FMRFamide relations).
                                                                                                                                                                                                                                                                                                                     Paemen L., Tips A., Schoofs L., Proost P., van Damme J., de Loof A.; "Lom-AG-myotropin: a novel myotropic peptide from the male accessory glands of Locusta migratoria."; Peptides 12:7-10(1991).

-!- FUNCTION: The exact physiological function is still unknown. This myotropic peptide is active on the oviduct and, to a lesser extent, on the hindgut. Transferred from the male to the female
                                                                                                                                                                                                  Neuropeptide; Amidation.
MOD_RES 13 13
SEQUENCE 13 AA; 1368 N
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MEDLINE=91271087; PubMed=2052501;
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NCBI_TaxID=7004;
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FMRFamide-like neuropeptide FLP7 (GYRKPPFNGSIF-amide).
Penaeus monodon (Penoeid shrimp).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE,
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TISSUE SPECIFICITY: Male
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Similarity 37.5%;
3; Conservative
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75.0%;
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RESULT 5
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FAR7 MACRS
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COXG_RAT
Neuropeptide;
MOD_RES 1
SEQUENCE 10
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NON TER
SEQUENCE
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01-NOV-1995
28-FEB-2003
                                                                                                                                                          Schaegger H., Noack H., Halangk W., Brandt U., von Jagow G.; "Cytochrome-c oxidase in developing rat heart. Enzymic properties amino-terminal sequences suggest identity of the fetal heart and tadult liver isoform.";
                                                                                                                                                                                                                                                                                                                                                                                        -!- SIMILARITY:
PIR; S65381; S6
                                                                                                                                                                                                                                                                                                                                                                                                                Eur. J. Biochem. 230:235-241(1995).

-!- FUNCTION: This protein is one of the nuclear-coded polypeptide chains of cytochrome c oxidase, the terminal oxidase in mitochondrial electron transport. This protein may be one of the heme-binding subunits of the oxidase.

-!- CATALYTIC ACTIVITY: 4 ferrocytochrome c + O(2) = 4 ferricytoch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=Wistar; TISSUE=Liver; MEDLINE=95324529; PubMed=7601105;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chor
                                                                                                                           SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                      Palaemonoidea;
                                                                                                                                                                                                                                                                                                                                                                               Oxidoreductase; Mitochondrion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cytochrome
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                          GO; GO:0007218;
                                                                                                                                           NCBI_TaxID=79674;
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(Rel. 32, Last sequence up
(Rel. 41, Last annotation
oxidase polypeptide VIb (
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                  Amidation.
                                                                                                                                                      Palaemonidae;
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                           P:neuropeptide signaling
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  1244
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Rodentia;
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100.0%;
  MW;
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 AMIDATION.
3CFE9C29C4540AA8 CRC64
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Sciurognathi; Muridae;
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                          pathway; IDA
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; Murinae; Rattus.
                                                                       rosembergii.";
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RESULT 6
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Eukaryota; Metazoa; Annelida; Clitellata; Oligochaeta; Hapl
Lumbricina; Lumbricidae; Eisenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1995 (Rel. 32, Last sequence up 01-NOV-1995 (Rel. 32, Last annotation Myoactive tetradecapeptide (ETP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sec
01-NOV-1995 (Rel. 32, Last and
         Holman G.M., Cook B.J., Nachman R.J.;
"Isolation, primary structure, and synthesis of leucoki
myotropic peptides of Leucophaea maderae.";
Comp. Biochem. Physiol. 88C:27-30(1987).
-!- FUNCTION: This cephalomyotropic peptide stimulates
activity of cockroach protodeum (hindgut).
-!- SUBCELLULAR LOCATION: Secreted.
PIR; JS0316; JS0316.
                                                                                                                                                                                                                                                                                       LEUMA
                                                                                                                                                                                                                                                                                                                                                                                                                                  Neuropeptide; Amidation.
MOD_RES 14 14
SEQUENCE 14 AA; 1478 |
                                                                                                                                                                            Leucophaea maderae (Madeira cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria;
                                                                                                                                                                                                               01-FEB-1991 (Rel. 17, Created)
01-FEB-1994 (Rel. 28, Last seq
10-OCT-2003 (Rel. 42, Last ann
Leucokinin VI (L-VI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptides 16:995-999(1995).
                                                                                                                                                                                                                                                   P19988;
01-FEB-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FISSUE=Gut;
MEDLINE=96087879; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nomoto K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=6396;
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                                                                                                       MEDLINE=87052651; PubMed=2877794;
                                                                                                                                                                  3laberidae; Leucophaea.
                                                                                                                     'ISSUE=Head;
                                                                                                                                EQUENCE
                                                                                                                                                      CBI_TaxID=6988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EQUENCE, AND SYNTHESIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SIMILARITY: TO INSECTS ALLATOTROPIN
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30.0%;
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Pyrrolidone
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annotation update)
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Pred. No. 7.3e

1; Mismatches
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Pred. No.
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CC9ABEF941CD91AD
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carboxylic acid
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                                                                                  leucokinins
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3laberoidea;
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Best Local
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MOD_RES
SEQUENCE
                                                                                                                                                                                                                           CXL1 CONMR
P58807;
28-FEB-2003
28-FEB-2003

    -!- FUNCTION: Degrades acharan sulfate and, heparin and heparan sulfate.
    -!- ENZYME REGULATION: Inhibited by cupric: Activated by reducing agents, such as DI

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=21223019; PubMed=11322884; Kim B.-T., Hong S.-W., Kim W.-S., Kim Y.S., Ki "Purification and characterization of acharan novel heparinases, from Bacteroides stercoris Eur. J. Biochem. 268:2635-2641(2001).
                                                                                                   Conus marmoreus (Marble cone).
Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;
Apogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NON_TER
                                                                                                                                                                                28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last seq
28-FEB-2003 (Rel. 41, Last ann
Lambda-conotoxin CMrVIA.
                                                                                                                                                                                                                                                                                                              CONMR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lyase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN-HJ-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteria; Bacteroidetes; Bac
Bacteroidaceae; Bacteroides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P83146;
28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence up
28-FEB-2003 (Rel. 41, Last annotation
Acharan sulfate lyase 1 (EC 4.2.2.-) (
MEDLINE=20564325; PubMed=10988292;
                    SEQUENCE, SYNTHESIS, TISSUE=Venom;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=46506;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacteroides stercoris.
                                                                              NCBI_TaxID=42752;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUBUNIT: Monomer.

PTM: The N-terminus is blocked.

MISCELLANEOUS: Has an isoelectric

7.2 and optimum temperature 45 deg

se; Heparin-binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mercaptoethanol.
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3; Conserv
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75.0%;
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                                          AND MASS SPECTROMETRY
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                                                                                                                                                                                                         sequence up
annotation
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Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01B2DAA241E865AB CRC64;
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is HJ-15.";
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DISULFID 3
MOD RES 10
SEQUENCE 11 AA; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                P58810;
28-FEB-2003
28-FEB-2003
28-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Lambda-conotoxins, a new family of conotoxins with unique disulfide pattern and protein folding. Isolation and characterization from the venom of Conus marmoreus.";

J. Biol. Chem. 275:39516-39522(2000).

-!- FUNCTION: Inhibits the neuronal noradrenaline transporter.

-!- SUBCELLULAR LOCATION: Secreted.

-!- TISSUE SPECIFICITY: Expressed by the venom duct.

-!- MASS SPECTROMETRY: MW=1237.93; MW_ERR=0.21; METHOD=Electrospray.

-!- SIMILARITY: Belongs to the chi/lambda-conotoxin family.
                                                                                         SEQUENCE
                                                                                                                                                                                                                                               Sharpe I.A., Gehrmann J., Loughnan M.L., Atkins A., Palant E., Craik D.J., Adams
                                                                                                                           DISULFID
                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Mollusca; Gastropoda; Orthogastropoda;
Apogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
NCBI_TaxID=42752;
                                                                                                                                                                                                                                                                                                                                                                                                                      28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Lambda/chi-conotoxin MrIB (Chi-MrIB).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Balaji R.A.,
Seow K.T., B
                                                                                                                                       Neurotoxin; Toxin;
                                                                                                                                                                                                                                                                                                                 MEDLINE=21419681; PubMed=11528421;
                                                                                                                                                                                                                                                                                                                                                                                                          Conus marmoreus (Marble cone).
                                                                                                                                                                                                                                                                                                                                        SEQUENCE, SYNTHESIS,
                                                                                                                                                                                                         . Neurosci. 4:902-907(2001).
FUNCTION: Inhibits the neuronal noradrenaline transporter.
SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expressed by the venom duct.
                                                                                                                                                            SIMILARITY: Belongs to
                                                                                                                                                                                              PTM: Exists in
                                                                                                                                                     11-Hyp-12.
MASS SPECTROMETRY: MW=1393.52; METHOD=Electrospray.
CTMTLARITY: Belongs to the chi/lambda-conotoxin family.
                                                                                                                                                1IEO; 03-APR-02
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b
                                            Similarity 3; Conserv
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                                             Conservative
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œ
                                                                                         AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                         1382
                                                                                                                                     Hydroxylation; 3D-structure
                                                                                                                                                                                                two forms,
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                                       30.0%; oc
100.0%; Pr
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                                                                                                                                                                                               Expressed : forms, due
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                                           Score 18; DB; Pred. No. 1.5
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                                                                                        HYDROXYLATION.
277AAC376EAD2B58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYDROXYLATION.
277AAC60B7232B58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                by the venom
to cis-trans
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o. 1.3e+03;
o;
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                                                                   DB 1;
                                                      1.5e+03;
                                                                                                                                                                                                                                                                                         Thomas L., Ac
D.J., Alewood
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                                                                                                                                                                                                                                                                   alphal-adrenoceptor
                                                                                         CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CRC64;
                                                                  Length 13;
                                                                                                                                                                                                duct.
isomerization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 11;
                                             Indels
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od P.F.,
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on from the
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RESULT 11

Best Loc Matches

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Similarity 3; Conser

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COXM_RA
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Best Local S
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4_RAT

COXM_RAT

P80431;

01-NOV-1995

01-NOV-1995

28-FEB-2003
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01-CCT-1994
01-NOV-1995
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P38495;
                                                                                                    Eur. J. Biochem. 230:235-241(1995).

-!- FUNCTION: This protein is one of the nuclear-coded polypeptide chains of cytochrome c oxidase, the terminal oxidase in mitochondrial electron transport.

-!- CATALYTIC ACTIVITY: 4 ferrocytochrome c + O(2) = 4 ferricytochrome
                                                                                                                                                        MEDLINE=95324529; PubMed=7601105;
Schaegger H., Noack H., Halangk W., Brandt U., von Jagow G.;
"Cytochrome-c oxidase in developing rat heart. Enzymic properties
amino-terminal sequences suggest identity of the fetal heart and t
adult liver isoform.";
                                                                                                                                                                                                                                                 Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Callinectes sapidus (Blue crab).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyu
Eubrachyura; Portunoidea; Portunidae; Callinectes.
                                             PIR; S65387; S65387.
Oxidoreductase; Mitochondrion.
NON TER 10 10
SEQUENCE 10 AA; 1210 MW; CI
                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eubrachyura; Por
NCBI_TaxID=6763;
                                                                                                                                                                                                                                                                                                                        Cytochrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sapidus."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cardioactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FMRFamide-like
                                                                                                                                                                                                                     STRAIN=Wistar; TISSUE=Liver;
                                                                                                                                                                                                                                                                                                   COX7B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptides 12:1295-1302(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Krajniak K.G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=92270479;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "The identification and structure-activity relations cardioactive FMRFamide-related peptide from the blue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -!- SIMILARITY:
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                                                                                                                                                                                                                                                                                                                       (Rel. 32, Created)
(Rel. 32, Last sequence update)
(Rel. 41, Last annotation updat
coxidase polypeptide VIIb, mito
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(Rel. 30, Last sequence update)
(Rel. 32, Last annotation update)
  28.3%;
larity 50.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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Y: Belongs to
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 Score 17; DB Pred. No. 1.7e 0; Mismatches
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134F0729D5A4045B CRC64;
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Pred.
                                               CFC70EB771A33326
                                                                                                                                                                                                                                                              Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Rattus
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                                                                                                                                                                                                                                                                                                                         update)
mitochondrial
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                         DB 1;
..7e+03;
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                                               CRC64;
                        Length 10;
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RESULT 14

NXSN_PSETE

ID __NXSN_P
AC _P59072

DT _28-FEB
DT _28-F
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Best Local S
Matches 3
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28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
28-FEB-2003 (Rel. 41, Last annotation update)
Short neurotoxin N1 (Alpha neurotoxin) (Fragment).
Pseudonaja textilis (Eastern brown snake).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Elapidae; Acanthophiinae; Pseudonaja.
NCBI_TaxID=8673;
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Flengsrud R., Skjeldal L.;
"Two-dimensional gel electrophoresis separation and N-terminal
"Two-dimensional gel electrophoresis separation and N-terminal
sequence analysis of proteins from Clostridium pasteurianum W5.";
Electrophoresis 19:802-806 (1998).

-!- FUNCTION: This protein promotes the GTP-dependent translocation of the P-site of
                                                                                                      Gong N.L., Armugam A., Jeyaseelan K.;
"Postsynaptic short-chain neurotoxins from Pseudonaja textilis:
cloning, expression and protein characterization.";
Eur. J. Biochem. 265:982-989(1999).
-!- FUNCTION: Lethal neurotoxin, binds and inhibits nicotinic
acetylcholine receptors (nAChR).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Elongation factor; Protein biosynthesis; GTP-binding NON TER 11 11 11 SEQUENCE 11 AA; 1337 MW; 412E71F1D9C33B17 CRC64;
                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE, AND TISSUE=Venom;
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Clostridium
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15-JUL-1998 (Rel. 36, Last seq
28-FEB-2003 (Rel. 41, Last ann
Elongation factor G (EF-G) (CP
                                                                                                                                                                                                                                                                                                                                           MEDLINE=99449602; PubMed=10518793;
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erPro; IPR000795; EF GTPbind.
SITE; PS00301; EFACTOR GTP; PARTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SUBCELLULAR LOCATION: Cytoplasmic. SIMILARITY: Belongs to the GTP-binding
                                  SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expressed by the venom gland
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Firmicutes; Clc
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   MW=6236; METHOD=Electrospray.
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1; Mismatches
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RESULT 16
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                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  UNSURE
NON_TER
SEQUENCE
                                                Eukaryota; Metazoa; Arthro
Neoptera; Orthopteroidea;
Blaberidae; Leucophaea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
01-NOV-1995 (Rel. 32, Last annotation update)
Myoactive tetradecapeptide (PTP).
Pheretima vittata (Earthworm).
Pheretima vittata (Earthworm).
Eukaryota; Metazoa; Annelida; Clitellata; Oligochaeta;
Lumbricina; Megascolecidae; Pheretima.
NCBI TaxID=46674;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PHEVI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SIMILARITY: Belongs to the snake toxin f InterPro; IPR003571; Snake toxin. PROSITE; PS00272; SNAKE_TOXIN; PARTIAL. Toxin; Neurotoxin; Postsynaptic neurotoxin; Acetylcholine receptor inhibitor; Multigene
                                                                                             01-MAY-1991 (Rel. 18, Created)
01-MAY-1991 (Rel. 18, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
Leucokinin IV (L-IV)
                                                                                                                                                                                                                                                                                                 Neuropeptide; Amidation.
MOD_RES 14 14
SEQUENCE 14 AA; 1522 MW;
                                                                                                                                                                                                                                                                                                                                 muscles.
-!- SIMILARITY: TO INSECTS ALLATOTROPIN
                                                                                                                                                                                                                                                                                                                                                                  Peptides 16:995-999(1995)
                                                                                                                                                                                                                                                                                                                                                                               "A novel
foetida."
                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=96087879; PubMed=8532604; Ukena K., Oumi T., Matsushima O.,
                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE, AND TISSUE=Gut;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P46980;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -!- MISCELLANEOUS: LD(50) is 0.84 mg/kg by -!- SIMILARITY: Belongs to the snake toxin
                                        VCBI_TaxID=6988;
                                                                                                                                                                                                                                                                                                                                                                                                     Nomoto
                                                                                     eucophaea maderae (Madeira cockroach)
                                                                                                                                                                                                                                                                                                                                                       FUNCTION: Has a stimulative effect on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PHEVI
                                                                                                                                                      LEUMA
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2; Conserv
                                                                                                                                                                                                                                                      Similarity 2; Conserv
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                                                                                                                                                                                                            GFRDGSADRI 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GYKNS 5
                                                                                                                                                                                                                                GYKNSKFHRV 10
                                                                                                                                                                                                                                                                                                                                                                                       gut tetradecapeptide isolated from
                                                                            Metazoa; Arthropoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               SYNTHESIS
                                                                                                                                                      STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1319 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28.3%;
40.0%;
                                                                                                                                                                                                                                                                 28.3%;
20.0%;
                                                              Dictyoptera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 17; DB
Pred. No. 1.9e
3; Mismatches
                                                                                                                                                                                                                                                      Score 17; DB 1;
Pred. No. 2.5e+03
4; Mismatches '
                                                                                                                                                                                                                                                                                                  AMIDATION.
DA40BEE67CCD91AD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0D1EF0C81B58732B CRC64;
                                                               Hexapoda; Insecta;
ptera; Blattaria;
                                                                                                                                                                                                                                                                                                                                                                                                              Ikeda
                                                                                                          update)
on update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.9e+03
                                                                                                                                                      ထ
                                                                                                                                                                                                                                                                                                                                                                                                           7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                      AA.
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                                                                                                                                                                                                                                                                                                                                                       the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             intravenous family.
                                                                                                                                                                                                                                                                                                                                                                                          the
                                                                                                                                                                                                                                                                                                                                                                                                             Fujita T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   family.
                                                                                                                                                                                                                                                                                                   CRC64
                                                                                                                                                                                                                                                                                                                                                         contraction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 11;
                                                                                                                                                                                                                                                                            Length 14
                                                                                                                                                                                                                                                                                                                                                                                          earthworm,
                                                                                                                                                                                                                                                                                                                                                                                                              Minakata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Haplotaxida;
                                                                                                                                                                                                                                                                                                                                                        O.f
                                                                                                                                                                                                                                                                                                                                                                                          Eisenia
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Holman

G.M., Cook B.J., Nachman

R.J.;

SEQUENCE, ANTISSUE=Head;

AND SYNTHESIS

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RESULT
FAR2 PE
AC PR
AC P
AC PR
AC P
  RESULT 18
TKL4_LOCMI
ID TKL4
AC P3025
DT 01-AF
DT 10-OC
DT LOCUS
OC LOCUS
OC EUKAZ
OC Neopt
OC NCBI_
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Best Local S
Matches 3
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Best Local S
Matches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuropeptide; 10 MOD_RES 10 MOD_R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sithigorngul P., Pupuem J., Krungkasem C., Longyant Chaivisuthangkura P., Sithigorngul W., Petsom A.; "Seven novel FMRFamide-like neuropeptide sequences f of the giant tiger prawn Penaeus monodon."; Comp. Biochem. Physiol. 131B:325-337(2002).
-!- SUBCELULAR LOCATION: Secreted.
-!- SUBCELULAR LOCATION: MW=1260.0; METHOD=MALDI.
                                                                                                                         01-APR-1993 (Rel. 25, Created)
01-APR-1993 (Rel. 25, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation updat
Locustatachykinin IV (TK-IV).
Locusta migratoria (Migratory locust).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Comp.
                                     Neoptera; Orthopteroidea;
Acridoidea; Acrididae; Oe
                                                                                                   Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Penaeus monodon (Penoeid shrimp).
Eukaryota; Metazoa; Arthropoda; Crus
Eumalacostraca; Eucarida; Decapoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
FMRFamide-like neuropeptide FLP2 (AYSNLNYLRF-amide).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FAR2_PENMO
P83317;
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MOD_RES 8 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Primary structure and synthesis of from Leucophaea maderae: members of Cephalomyotropins.";
                                                                                                                                                                                                                                                                   P30250;
01-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GO; GO:0007218; P:neuropeptide signaling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=21956277; PubMed=11959015;
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                                                                                                                                                                                                                                                                                                                                    TKL4_LOCMI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=6687;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Penaeidae; Penaeus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Penaeus monodon (Penoeid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FUNCTION: This cephalomyotropic activity of cockroach protodeum SUBCELLULAR LOCATION: Secreted.
_TaxID=7004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   family.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YKNSKFHR
                                   Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Orthopteroidea; Orthoptera; Caelifera; Acridomorpha
; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 AA; 906 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10 AA;
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                                                                                                                                                                                                                                                                                                                                      STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1260 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26.7%;
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Pred. No. 1.4e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 16;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Crustacea; Malacostraca;
oda; Dendrobranchiata; Penaeoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  two additional neuropeptides a new family of
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                                                                                                                                                                                                     update)
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RESULT 19
ANGT HORSE
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Query Match
Best Local S
Matches 2
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Best Local Similarity
Matches 3; Conser
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-ANGT_HORSE

P01016;

21-JUL-1986

21-JUL-1986

10-OCT-2003
                                                                                                                                                      Vasoconstrictor; Plasma; S
PEPTIDE 1 10
PEPTIDE 1 8
PEPTIDE 2 8
NON TER 14 14
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SEQUENCE.
TISSUE=Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                   Skeggs L.T. Jr., Kahn J.R., Lentz K., Shumway N.P.;
"The preparation, purification, and amino acid sequence of a polypeptide renin substrate.";
J. Exp. Med. 106:439-453(1957).

-!- FUNCTION: In response to lowered blood pressure, the enzoleaves angiotensin I, from angiotensinogen. ACB (angiotensverting enzyme) then removes a dipeptide to yield the physiologically active peptide angiotensin II, the most physiologically active peptide angiotensin II, the most
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Regul.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria;
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Equus caballus (Horse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oviduct and foregut.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: SOME SIMILARITY TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             with homology to peptides of Regul. Pept. 31:199-212(1990)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Angiotensinogen II); Angiotensin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Locustatachykinin III and IV: two additional insect
                                                                                                                                                                                                                                                                                                                   InterPro; IPR0002: PROSITE; PS00284;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGT OR SERPINA8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   achykinin; Neuropeptide; Amidation
                                                                                                                                                                                                                                                                                                                   ; 1ER8; 15-UL
PR000215; S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: Myoactive peptide.
                                                                                                                                                                                                                                                                                                                                                                                                              A92775; A01250
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  Similarity
2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (Rel. 01, Created)
(Rel. 01, Last sequence update)
(Rel. 01, Last sequence update)
(Rel. 42, Last annotation update)
ogen [Contains: Angiotensin I (Ang I); Angiotensin II (Angiotensin II)]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
     Conservative
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                                                                                                                                                                     1-91.
215; Serpin.
215; Serpin.
3; SERPIN; PARTIAL.
4; Plasma; Serpin; 3D-structure.
5; Plasma; Serpin; 3D-structure.
10 ANGIOTENSIN II.
10 ANGIOTENSIN III.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                     IFICITY: Expressed by the liver and Belongs to the serpin family.
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                                                                                                                                 1759
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                                    26.7%;
40.0%;
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75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 16; DB:
Pred. No. 2.7e-
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        o
;-
       Score 16; DB Pred. No. 3.9e 2; Mismatches
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                                                                   DB 1;
                                         9e+03;
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                                                                                                                                   CRC64;
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                                                                   Length
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          Indels
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RS19 CLOPP
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Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                              RS19_LOWBP
Q48878;
30-MAY-2000
                                                                                                                                                                                                    RPSS OR RPS19.

Loofah witches'-broom phytoplasma.

Loofah wirmicutes; Mollicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                LT 21
LOWBP
RS19_I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q46228;
30-MAY-2000
30-MAY-2000
28-FEB-2003
                SEQUENCE FROM N.A.
MEDLINE=94350802; PubMed=8071198;
Gundersen D.E., Lee I.M., Rehner S.A.,
"Phylogeny of mycoplasmalike organisms
their classification.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                               30-MAY-2000
28-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HAMAP; MF_00531; -; 1.
InterPro; IPR002222; Ribosomal_S19.
PROSITE; PS00323; RIBOSOMAL_S19; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=94350802; PubMed=8071198;
Gundersen D.E., Lee I.M., Rehner S.A.,
"Phylogeny of mycoplasmalike organisms
their classification.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ribosomal protein; rRNA-binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; L27011; AAA83938.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Clover proliferation phytoplasma.
Ranteria: Firmicutes; Mollicutes; Acholeplasmatales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                           Acholeplasmataceae;
NCBI_TaxID=35773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=35776;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Acholeplasmataceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RPSS OR RPS19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0S ribosomal protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bacteriol. 176:5244-5254(1994). - FUNCTION: Protein S19 forms a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to the 16S ribosomal RNA (By similarity).
SIMILARITY: Belongs to the S19P family of ribosomal proteins.
                                                                                                                                                                                                                                                                                         ribosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLOPP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 KNSK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 3; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KNOK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14 AA;
                                                                                                                                                                                                                                                                                                           (Rel. 39, Created)
(Rel. 39, Last seg
(Rel. 41, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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(Rel. 39, Last sequence up
(Rel. 41, Last annotation
al protein S19 (Fragment).
                                                                                                                                                                                                                                                                                     protein
                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1642 MW;
                                                                                                                                                                                    Phytoplasma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Phytoplasma.
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75.0%;
                                                                                                                                                                                                                                                                                       Last
S19
                                                                                                                                                                                                                                                                                                         Last sequence update)
                                                                                                                                                                                                                                                                                       (Fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20C478EB9FFFE4A8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                Acholeplasmatales;
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                                      Davis R.E., Kingsbury D.T.; (phytoplasmas): a basis for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Davis R.E., Kingsbury D.T.; (phytoplasmas): a basis for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; 1
3.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                            AA
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Matches 3
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P37985;
01-OCT-1994
01-OCT-1994
01-NOV-1995
                                                                                                                                                                                                                                                                                              Douillie A., Toussaint A., Faelen M.;
Submitted (AUG-1993) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: BINDS MRNA; THUS FACILITATING RECOGNITION OF THE
INITIATION POINT. IT IS NEEDED TO TRANSLATE MRNA WITH A SHOR
SHINE-DALGARNO (SD) PURINE-RICH SEQUENCE (BY SIMILARITY).
-!- SIMILARITY: Belongs to the S1P family of ribosomal proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ribosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HAMAP; MF 00531; -; 1.
HAMAP; MF 00531; -; 1.
InterPro; IPR002222; Ribosomal S19.
PROSITE; PS00323; RIBOSOMAL S19; PARTIAL.
Ribosomal protein; rRNA-binding.
                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
                                                                                            SEQUENCE
                                                                                                        Ribosomal protein; Repeat; RNA-binding.
                                                                                                                                                    EMBL; X74750; CAA52769.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=3937;
                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Pectobacterium.
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                                                                                                                                       S37141; S37141.
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(Rel. 30, Last seq
(Rel. 32, Last ann
                                 Conservative
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, Last annotation on
n S1 (Fragment).
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75.0%;
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Pred. No. 1.4e
3; Mismatches
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Pred. No.
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me S19P family of ribosomal
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3.9e+03;
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Hughes G.J., Frutiger S., Paquet N., Pasquali C., S.
Tissot J.-D., Bairoch A., Appel R.D., Hochstrasser
"Human liver protein map: update 1993.";
Electrophoresis 14:1216-1222(1993).
-!- MISCELLANEOUS: On the 2D-gel the determined pI
protein is: 6, its MW is: 15 kDa.
SWISS-2DPAGE; P31929; HUMAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-AUG-1988
01-AUG-1988
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Intestine;
MEDLINE=86192829; PubMed=2422058;
Conlon J.M., Deacon C.F., O'Toole
"Scyliorhinin I and II: two novel
FEBS Lett. 200:111-116(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P31929;
01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of liver tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SCYCA
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
NCBI_TaxID=9606;
                                                                 muscles.
-i- SUBCELLULAR LOCATION: S-i- SUBCELLULAR LOCATION: S-i- SUBCELLULAR ITY: Belongs to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyliorhinidae; Scyliorhinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TKON1_SC
                                                                                                                                                                                                                                                                                         Waugh D., Wang Y., Hazon N., Balment R.J., Conlon J.M.; "Primary structures and biological activities of substance-P-related peptides from the brain of the dogfish, Scyliorhinus canicula."; Eur. J. Biochem. 214:469-474(1993).
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                                                                                                                                                                                    evoke behavioral responses, are potent vasodilators secretagogues, and contract (directly or indirectly)
                                                                                                                                                                                                                                                        FUNCTION: Tachykinins are active peptides which excite
                                                                                                                                                                                 secretagogues,
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iproo2040; Tachy_Neurokinin
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tachykinins
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RESULT 26
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P82092;
16-OCT-2001
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"Caerulein-like peptides from the skin glands o
montains tree frog Litoria citropa. Part 1. Seq
using electrospray mass spectrometry.";
Rapid Commun. Mass Spectrom. 13:2498-2502(1999)
-!- FUNCTION: Hypotensive neuropeptide (Probable)
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin dorsal glands.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00267; TACHYKININ; 1.
Tachykinin; Neuropeptide; Amidation.
MOD_RES 10 10 AMIDATI
SEQUENCE 10 AA; 1219 MW; D0602D0
                                                                                                    TKC2_CAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Litoria citropa (Australian blue mountains tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
Calliphora vomitoria (Blue blowfly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oe
Calliphoridae; Calliphora.
                                                             01-NOV-1995 (Rel.
01-NOV-1995 (Rel.
10-OCT-2003 (Rel.
                                                                                                                                                                                                                                                                                                                                        InterPro; IPR001651; Gastrin.
PROSITE; PS00259; GASTRIN; FALSE_NEG.
Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation;
                                                                                                                                                                                                                                                                                                                                                                                                                      -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation updat
Caerulein 4.2/4.2Y4.
                                                    Callitachykinin II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=20057701; PubMed=10589099;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=94770;
                                                                                                                                                                                                                                                                                                                                                                                  -!- MASS SPECTROMETRY: MW=1404; METHOD=Electrospray.
-!- SIMILARITY: Belongs to the gastrin/cholecystokinin family.
                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE SPECIFICITY: Skin dorsal PTM: Isoform 4.2Y4 differs from
                                                                                                                 CALVO
                                                                                                                                                                                                                                                                                                                                   idone carboxylic acid.
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3; Conserv
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                                                                                                                                                                                                       YKNSKF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                   11 AA;
                                                            (Rel. 32, Created)
(Rel. 32, Last sec
(Rel. 42, Last ann
                                                                                                                                                                                                                                  Conservative
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                                                                                                                 STANDARD;
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                                                                                                                                                                                                                                                                                    1344 MW;
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50.0%;
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50.0%;
                                                               sequence up
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1; Mismatches.
                                                                                                                                                                                                                                 Score 15; DB Pred. No. 4.6e 0; Mismatches
                                                                                                                                                                                                                                                                                              PYRROLIDONE SULFATION. AMIDATION.
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D0602D6B59C33AA9
                                                                                                                                                                                                                                                                                    10DAB894F5B861BB
                                                                                                                  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                           glands.
n isoform
                                                                          update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     glands of the rt 1. Sequence
                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                       CARBOXYLIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B
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                                                                                                                                                                                                                                               ,6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                             4.2 in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 10;
                                                                                                                                                                                                                                                          Length 11;
                                                                                                                                                                                                                                   Indels
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                Oestroidea;
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RESULT

NP1_LYM

AC PR

AC PR

DT 01

ST 01

     RESULT

NP4_LYM

NP4_LYM

AC P8

AC P8

DT 01

DT 01

DT 01

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DT 1
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Best Local S
Matches 3
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Best Local S
Matches 3
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NP1 LYMST
P80178;
                                                                                                                                                                                                                                                                                                 LYMST
                                                                                                                             NP4 LYMST
P80181;
01-JUL-1993
01-JUL-1993
01-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOD RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Ganglion;
MEDLINE=93238777; PubMed=8477756;
Johnsen A.H., Rehfeld J.F.;
"LymnaDFamides, a new family of neuropeptides from the pond Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
Lymna-DF-amide 1.
Lymnaea stagnalis (Great pond snail).
Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata;
Lymnaeoidea; Lymnaeidae; Lymnaea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lymnaea stagnalis (Great pond snail).
Eukaryota; Metazoa; Mollusca; Gastrop
Lymnaeoidea; Lymnaeidae; Lymnaea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tachykinin; Neuropeptide; Amidation.

MOD_RES 11 11 AMIDATION.

SEQUENCE 11 AA; 1103 MW; 15D7E3F9C9CDD444 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE, AND SYNTHESIS.

MEDLINE=95075727; PubMed=7984492;

Lundquist C.T., Clottens F.L., Holman G.M., Nichols
Naessel D.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Neuropeptide; Amidation.
MOD_RES 13 13
SEQUENCE 13 AA; 1519 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Callitachykinin I and II, two novel the blowfly, Calliphora vomitoria, th tachykinins."; Peptides 15:761-768(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=27454;
[1]
SEQUENCE, AND SYN
                                                                                                   Lymna-DF-amide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nvertebrates?";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . J. Biochem. 213:875-87
SIMILARITY: RELATED TO
; S32471; S32471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FUNCTION: Myoactive peptide.
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: SOME SIMILARITY TO TACHYKININS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              __TaxID=6523;
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3; Conserv
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(Rel.
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                                                                                                                                                                                                                                                                STANDARD;
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                                                                                                                        . 26, Created)
. 26, Last sequence.
. 26, Last anno
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RELATED TO THE CHOLECYSTOKININ
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75.0%;
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42.9%;
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:
                                                                                                                             sequence update) annotation updat
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Pred. No. 4.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AMIDATION.
9CA07BA3F5D5B455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gastropoda; Pulmonata; Basommatophora;
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                                                                                                                                update)
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;
                                  Basommatophora;
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RESULT
RS19_PF
ID RS
AC QS
AC QS
DT 30
DT 30
DT 30
DT 28
GN RF

PPWBP STANDARD; PRT; 14 AA. Q52093; 30-MAY-2000 (Rel. 39, Created) 30-MAY-2000 (Rel. 39, Last sequence update) 28-FEB-2003 (Rel. 41, Last annotation update) 30S ribosomal protein S19 (Fragment). RPSS OR RPS19.

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RESULT
TEML_RA
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Best Local S
Matches 3
                                                                              Matches
                                                                                                  Query Match
Best Local :
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TEML RANTE
P57104;
16-OCT-2001
16-OCT-2001
                                                                                                                                       Gram-positive Contion: Secreted.
-!- SUBCELULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin.
-!- SIMILARITY: Belongs to the brevinin family.
-!- SIMILARITY: Belongs to the brevinin family.
Amphibian defense peptide; Antibiotic; Amidation.
MOD RES
13
13
AMIDATION.
13 AA; 1641 MW; 9EBDCB1FAFF7C325 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eur. J. Blocken. -
-!- SIMILARITY: RELATED
PIR; S32474; S32474.

Pir: S32474; S32474.
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SEQUEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rana temporaria (European com
Eukaryota; Metazoa; Chordata;
Amphibia; Batrachia; Anura; Ne
NCBI_TaxID=8407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuropeptide; Amidation.
MOD_RES 13 13
SEQUENCE 13 AA; 1503 |
                                                                                                                                                                                                                                                                                                    Eur. J. Bioche
                                                                                                                                                                                                                                                                                                                                         "Temporins,
temporaria."
                                                                                                                                                                                                                                                                                                                                                                                   Barra D.
                                                                                                                                                                                                                                                                                                                                                                                                    Simmaco M.,
                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=97175050; PubMed=9022710;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=6523;
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                                                                                                                                                                                                                                                                             FUNCTION: Has antibacterial Gram-positive bacteria.
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3; Conserv
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                                                                                                 Similarity
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40, Last sequence up
42, Last annotation
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Anura; Neobatrachia; Ranoidea; Ranidae; Rana
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9CA07BBB56D5B455
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NCBI_TaxID=8097, 8019, 7936;
                             SPECIES=O.kisutch; TISSUE=Pancreas;
MEDLINE=87055212; PubMed=2877919;
Plisetskaya E.M., Pollock H.G., Rouse J.B., Hamilton J.W.,
Kimmel J.R., Andrews P.C., Gorbman A.;
"Characterization of coho salmon (Oncorhynchus kisutch) islet
                                                                                                                                                                                                                                                                                                                     SPECIES=M.scorpius; TISSUE=Pancreas;
MEDLINE=88029486; PubMed=2889597;
Conlon J.M., Davis M.S., Falkmer S., Thim L.;
"Structural characterization of peptides derived from prosomatostatins I and II isolated from the pancreatic islets of prosomatostatins I and II isolated from the pancreatic islets of species of teleostean fish: the daddy sculpin and the flounder.";
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    -!- FUNCTION: Protein S19 forms a complex with S13 that binds stito the 16S ribosomal RNA (By similarity).
    -!- SIMILARITY: Belongs to the S19P family of ribosomal proteins.

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-!- FUNCTION: Protein S19 forms a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=94350802; PubMed=8071198;
Gundersen D.E., Lee I.M., Rehner S.A.,
"Phylogeny of mycoplasmalike organisms"
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Bacteria; Firmicutes
Acholeplasmataceae;
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InterPro; IPR002222; Ribosomal_S19.
PROSITE; PS00323; RIBOSOMAL_S19; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Myoxocephalus scorpius (Shorthorn sculpin) (Daddy sculpin), Oncorhynchus kisutch (Coho salmon), and Anguilla anguilla (European freshwater eel).

Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Eutelec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement (S or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This
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J. Biochem.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     L27036; AAA83946.1; -.
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SEQUENCE
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SPECIES=A.mississippiensis; TISSUE=Stomach; SPECIES=3324451; PubMed=8101369;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SPECIES=A.anguilla; TISSUE=Pancreas;
MEDLINE=89065329; PubMed=2904391;
Conlon J.M., Deacon C.F., Hazon N., Henderson I.W.,
"Somatostatin-related and glucagon-related peptides
structural features from the European eel (Anguilla
Gen. Comp. Endocrinol. 72:181-189(1988).
-i- FUNCTION: Somatostatin inhibits the release of s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gen.
                                                                                                                                                                                                                                                                                                                                                                                                 "Isolation and structural characterization of insulin, "Isolation and structural characterization of insulin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hormone, Multigene family.
DISULFID 3 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES=A.anguilla;
                                                                                                                                                                                                                                                                                                                               Peptides 11:461-466(1990).
-!- FUNCTION: Somatostatin inhibits
-!- SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES=T.scripta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metāzoa; Chordata; Craniata; Vertebrata;
Archosauria; Crocodylidae; Alligatorinae; Alligator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Somatostatin-14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Neuroendocrine peptides (NPY, and stomach of the alligator."; Peptides 14:573-579(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wang Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alligator mississippiensis (American alligator),
Trachemys scripta (Red-eared slider turtle) (Pse
                                                                                                                                                                                                                                                                       -!- SIMILARITY:
PIR; C60414; C60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=90341082;
                                                                                                                                                                                  Hormone
                                                                                                                                                                                                            Pfam; PF03002; Somatostatin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CBI_TaxID=8496,
                                                                                                                                                                                                                                            nterPro;
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:- SIMILARITY: Belongs to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B60842; B60842
S00172; S00172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      S00172;
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                                                                                                                                                                                                                                                                    C60414; C60414.
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Similarity 3; Conserv
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                                                                                                                                                                                                                                            IPR004250;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 AA;
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(Rel. 26, Last sequence up
(Rel. 40, Last annotation
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75.0%;
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ostatin; 1.
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Pred.
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                                                                                                                       D6270F5C09682679
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GRP,
                                                                                                                                                                                                                                                                                                    somatostatin
                                                                                                                                                       SIMILARITY.
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No.
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6e+03;
                                  DB 1;
6e+03;
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                                                                                                                         CRC64;
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with unu
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RESULT 34
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FAR4_CALVO
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Matches 3
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P41859;
01-NOV-1995
01-NOV-1995
01-NOV-1995
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SEQUENCE
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"Isolation, structure, and activity of -Phe-Met-Arg-Phe-NH2 neuropeptides (designated calliFMRFamides) from the blowfly Calliphora vomitoria.";
Proc. Natl. Acad. Sci. U.S.A. 89:2326-2330(1992).
                                                                                                                                      Nakamura S., Takenaka O., Takahashi K.;
"Fibrinopeptides A and B of baboons (Papio anubis, Pagand Theropithecus gelada): their amino acid sequences evolutionary rates and a molecular phylogeny for the b J. Biochem. 94:1973-1978(1983).

-!- FUNCTION: Fibrinogen has a double function: yieldipolymerize into fibrin and acting as a cofactor in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last seq
10-OCT-2003 (Rel. 42, Last ann
Fibrinogen beta chain [Contain
                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota;
Mammalia; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Calliphora vomitoria (Blue blowfly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Calliphoridae; Calliphora.
Calliphoridae; Calliphora.
                                                                                                                                                                                                                                                                                                                                                                       Cercopithecinae;
NCBI_TaxID=9557;
                                                                                                                                                                                                                                                                                                                                                                                                                 Papio hamadryas (Hamadryas baboon).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neuropeptide;
                                                                                                                                                                                                                                                                                                                                  SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Thoracic ganglion;
MEDLINE=92196111; PubMed=1549595;
Duve H., Johnsen A.H., Sewell J.C.,
                                                 --
                                                                                                                                                                                                                                                                                                    MEDLINE=84161822; PubMed=6423621;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CalliFMRFamide
                                                                                            aggregation.
SUBUNIT: HEX
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                   SUBUNIT: HEXAMER CONTAINING 2 SETS O (ALPHA, BETA AND GAMMA), LINKED TO E PTM: Conversion of fibrinogen to fib which cleaves fibrinopeptides A and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PAPHA
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3; Conserv
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9 9
9 AA; 1182 MW;
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(Rel. 32, Last seq
(Rel. 32, Last ann
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Last sequence update)
Last annotation update)
[Contains: Fibrinopeptide
                                                                                                                                        as a double function: yielding monomers and acting as a cofactor in platelet
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Pred. No. 1
0; Mismatch
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31730699CAB6D457
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3 NONIDENTICAL CHAINS
CH OTHER BY DISULFIDE BONDS.
in is triggered by thrombin,
in is triggered beta chains,
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                                                                                                                                                                                                                                                       Papio hamadryas,
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Query Match Best Local

Similarity

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Score Pred.

14; No.

DB 1; 1.4e+05;

Length

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Histidine metabolism; NON TER 9 SEQUENCE 9 AA; 114

1140 MW;

970FC41B5325A6C5 CRC64;

Lyase;

NAD.

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NON TER
SEQUENCE
                                                                                                                         between the Swiss Institute of Bioinformatics and the Ethe European Bioinformatics Institute. There are no rest use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an entities requires a license agreement (See http://www.isb-orsend an email to license@isb-sib.ch).
                                                                                                                                                                                                                                      yl)propanoate = urocanate + H(2)O.

-!- COFACTOR: Binds 1 NAD per subunit (By similarity).

-!- PATHWAY: Histidine degradation; second step.

-!- SUBCELLULAR LOCATION: Cytoplasmic (By similarity).

-!- SIMILARITY: Belongs to the urocanase family.
                                                                                                                                                                                                                                                                                                          Schwacha A., Bender R.A.;
"Nucleotide sequence of the gene encoding the repressor for institution utilization genes of Klebsiella aerogenes.";
J. Bacteriol. 172:5477-5481(1990).
-!- CATALYTIC ACTIVITY: 3-(5-0x0-4,5-dihydro-3-H-imidazol-4-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Klebsiella aerogenes.
Bacteria; Proteobacteria; Gámmaproteobacteria;
Proteobacteria; Klebsiella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PIR; E28854; E28854.
InterPro; IPR002181; Fibrinogen C.
PROSITE; PS00514; FIBRIN AG C DOMAIN; PARTIAL.
Blood coagulation; Plasma.
FIBRINOPEPTIDE B.
                                                    EMBL; M19665; AAA25078.1; -.
EMBL; M34604; AAA25076.1; -.
HAMAP; MF_00577; -; 1.
InterPro; IPR000193; Urocana
                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=88198018; PubMed=2834335; Nieuwkoop A.J.; Baldauf S.A., Hudspeth M.E.S., Bender R.A.; "Bidirectional promoter in the hut(P) region of the histidine utilization (hut) operons from Klebsiella aerogenes."; J. Bacteriol. 170:2240-2246(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-1989 (Rel. 12, Created)
01-OCT-1989 (Rel. 12, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Urocanate hydratase (EC 4.2.1.49) (Urocanase)
                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=90368611; PubMed=2203754;
                                            InterPro; IPR0001:
PROSITE; PS01233;
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E28854; E28854.
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1057
                                         93; Urocanase.
UROCANASE; PF
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RESIDENT TO SEQUENCE in a hypertensin.";

Princhem. J. 65:246-254(1957).

Biochem. J. 65:246-254(1957).

Cleaves angiotensin I, from angiotensinogen. ACE (angiotensive) then removes a dipeptide to yield the converting enzyme) then removes a dipeptide to yield the converting enzyme) then removes a dipeptide to yield the converting enzyme, which helps regulate volume and pressor substance known, which helps regulate volume and converting body fluids.

C is SUBCELLULAR LOCATION: Secreted.

C ISSUE SECIFICITY: Expressed by the liver and secreted in the secreted substance is a substance of body fluids.
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ANGT BOVIN STANDARD,
PO1017;
PO1017;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Angiotensinogen [Contains: Angiotensin I (Ang I); Angiotensin
Angiotensin III (Ang III) (Des-Asp[1]-angiotensin II)]
                                                                                                                                                 PVK_LOCMI
P83382;
28-FEB-2003
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PROSITE; PS00284; SERPIN; PARTIAL.

Vasoconstrictor; Plasma; Serpin; 3D-structure

PEPTIDE 1 0 ANGIOTENSIN II.

PEPTIDE 1 8 ANGIOTENSIN II.

PEPTIDE 2 8 ANGIOTENSIN III.

PON TER 10 10
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Elliott D
                                                                                                        28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation updat
Periviscerokinin (Lom-PVK-1).
   SEQUENCE, FUNCTION TISSUE=Abdominal
                                     Acridoidea; Acrididae;
NCBI_TaxID=7004;
                                                                   Locusta migratoria (Migratory locust).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Ac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bovidae; Bovinae;
NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea; Bovidae; Bovinae; Bos.
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  FUNCTION, AND MASS SPECTROMETRY dominal perisympathetic organs;
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100.0%; Pr
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P29221;
01-DEC-1992
01-DEC-1992
01-NOV-1995
                                                      InterPro; IPR000114; Ribosomal L16.
PROSITE; PS00586; RIBOSOMAL L16_1; PR0SITE; PS00701; RIBOSOMAL L16_2; Ribosomal protein; rRNA-binding.
NON TER 10 10
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Predel R., Gaede G.;
"Identification of the abundant neuropeptide from abdominal perisympathetic organs of locusts.";
Peptides 23:621-627(2002).
-i- FUNCTION: Myotropic peptide; increases the frequency of contraction of the heart and stimulates amplitude and to
                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sorsend an email to license@isb-sib.ch).
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MEDLINE=92210505; Publ
Lim P.O., Sears B.B.;
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MOD_RES 10 10
SEQUENCE 10 AA; 1105 N
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                                                                                                                                                                                                                                                                                                            gene sequences.
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Acholeplasma laidlawii.
Acholeplasma laidlawii.
Acholeplasma laidlawii.
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- FUNCTION: This protein binds located at the a site of the
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GO:0005184; F:neuropeptide hormone activity;
GO:0007218; P:neuropeptide signaling pathway
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MASS SPECTROMETRY: MW-
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Similarity 2; Conserv
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mal protein L16 (Frag
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                                          MEDLINE=66076612; PubMed=5857249;

Erspamer V., Anastasi A., Bertaccini G., Cei J.M.;

Erspamer V., Anastasi A., Bertaccini G., Cei J.M.;

"Structure and pharmacological actions of physalaemin, the main active polypeptide of the skin of Physalaemus fuscumaculatus.";

Experientia 20:489-490(1964).

-i- FUNCTION: Tachykinins are active peptides which excite neurons, evoke behavioral responses, are potent vasodilators and secretagogues, and contract (directly or indirectly) many smooth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P08615;
01-AUG-1988
01-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TRP8 LEUMA
P81740;
30-MAY-2000
30-MAY-2000
10-CCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tachykinin; Neuropeptide; Amidation.

MOD_RES 10 10 AMIDATI
SEQUENCE 10 AA; 1076 MW; 9E41037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptides 18:7-15(1997).

-!- FUNCTION: Myoactive peptide. Increases the of spontaneous contractions and tonus of hi -!- SUBCELLULAR LOCATION: Secreted.

-!- TISSUE SPECIFICITY: Brain.

-!- MASS SPECTROMETRY: MW=1076.9; METHOD=MALDI.

-!- SIMILARITY: SOME SIMILARITY TO TACHYKININS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IX
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30-MAY-2000 (Rel. 39, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Tachykinin-related peptide 8 (LemTRP 8).
Leucophaea maderae (Madeira cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blaberoidea;
Blaberidae; Leucophaea.
                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota;
Amphibia; E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Seven tachykinin-related peptides isolated from madeira cockroach; evidence for tissue-specific e
                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                    Leptodactylinae; Physalaemus.
NCBI_TaxID=8378;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=97269266; PubMed=9114447; Muren J.E., Naessel D.R.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         hysalaemus
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muscles.
SUBCELLULAR LOCATION: Secreted TISSUE SPECIFICITY: Skin.
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(Rel. 28, Last sequence update)
(Rel. 42, Last annotation update)
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Pred. No.
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9E410371E9C87685 CRC64;
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of hindgut i
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Leptodactylidae;
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01-JAN-1990
01-JAN-1990
10-OCT-2003
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16-OCT-2001
16-OCT-2001
28-FEB-2003
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MOD_RES
SEQUENCE
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Antibiotic: Bacterion:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -!- SIMILARITY: Belongs to the tachykinin family.
PIR; S07201; S07201.
InterPro; IPR002040; Tachy_Neurokinin.
Pfam; PF02202; Tachykinin; 1.
PROSITE; PS00267; TACHYKININ; 1.
Amphibian defense peptide; Tachykinin; Neuropeptide; Amidation;
Pyrrolidone carboxylic acid.
                                                                                                                                                                                                                                               Pattnaik P.;
"Biochemical characterization and molecular genetics of bacteriocin produced by a selected rumen bacterial isolate.";
Thesis (1999), National Dairy Research Institute / Karnal, India.
-!- FUNCTION: OXYGEN LABILE BACTERIOCIN-LIKE. HAS ANTIBACTERIAL ACTIVITY.
                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                              STRAIN=26L-10/3RA;
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NCE 12 AA; 1414 MW;
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11 AA;
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                            (Rel. 13, Created)
(Rel. 13, Last sequence update)
(Rel. 42, Last annotation update)
erase (EC 5.3.1.5) (Fragment).
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40, Last sequence up
41, Last annotation
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3293693E59C33457 CRC64;
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Streptomyces violaceoruber. Bacteria; Actinobacteria; Actinobacteridae;

Actinomycetales;

Xylose isomerase

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Best Local S
Matches 2
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P20304;

01-FEB-1991 (Rel. 17, Created)

01-FEB-1991 (Rel. 17, Last sequence update)

01-OCT-1996 (Rel. 34, Last annotation update)

Isocitrate dehydrogenase [NADP] cytoplasmic (EC 1.1.1.42)
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Biochem. J
                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=87308293; PubMed=2887570;
Bailey J.M., Colman R.F.;
"Isolation of the glutamyl peptide
2-(4-bromo-2,3-dioxobutylthio)-1,N(
2',5'-biphosphate in the active sit
dehydrogenase.";
                                                                                                                                              MEDLINE=87308292; PubMed=3624273;
Ehrlich R.S., Colman R.F.;
"Characterization of an active site position and active site position of an active site position of a color active site position of an active site position of active site pos
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InterPro; IPR001998; Xylose_isom.
PROSITE; PS00172; XYLOSE_ISOMERASE_1; PARTIAL PROSITE; PS00173; XYLOSE_ISOMERASE_2; PARTIAL PROSITE P
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-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the xylose
InterPro; IPR001804; Isodh.
PROSITE; PS00470; IDH_IMDH; PARTIAL
Oxidoreductase; NADP; Tricarboxylic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota;
Mammalia;
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-!- CATALYTIC ACTIVITY: D-xylose = D-
-!- COFACTOR: Binds 2 magnesium ions
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Vangrysperre W., Ampe C., Kerster
"Single active-site histidine in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=LMG 7183;
MEDLINE=90104230;
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NCBI_TaxID=1935;
                                                                                           -!- SUBUNIT: Homodimer.
                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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line in D-xylose isomerase from St
ion by chemical derivatization an
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P32560;
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28-FEB-2003
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ACT_SITE
NON_TER
SEQUENCE
 MEDLINE=21642653; PubMed=11756661;
Eigenheer R.A., Nicolson S.W., Schegg K.N
"Identification of a potent antidiuretic
Malpighian tubules.";
Proc. Natl. Acad. Sci. U.S.A. 99:84-89(20)
-!- FUNCTION: Strong inhibitor of fluid s
                                                                                                                                                                                                                                                                                                                                                                        ADFA_TEI
P82965;
                                                                                                                                                                                                       Tenebrio molitor (Yellow mealworm).

Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Coleoptera; Polyphaga; Cucujiformia;
Tenebrionidae; Tenebrio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Small intestine;
MEDLINE=86248085; PubMed=3087775;
Shaw C., Thim L., Conlon J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                    28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation updat
nntidiuretic factor A (ADFa) (ADF) (Antidiu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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MOD_RES 1 1 PYRROI
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                                                                                                                           FISSUE=Head;
                                                                                                                                                                                    VCBI_TaxID=7067;
                                                                                                                                                EQUENCE,
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!- FUNCTION: Smooth muscle-contracting peptide.
!- SUBCELLULAR LOCATION: Secreted.
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                                                                                                                                              FUNCTION,
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Izoa; Chordata;
Iria; Rodentia;
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Pred. No. 8.5e+03;
1; Mismatches 0;
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Pred. No. 8.5e+03;
0; Mismatches 3
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r of fluid secr
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Hystricognathi; Caviidae;
                                                                                                                                                                                                                                                                                         ion update)
(Antidiuretic
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                                             g K.M., ac
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                                                                                                                                                 SPECTROMETRY
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Copyright

GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                   Score
                                                                                                                                                                                                                                                                                                                                                                                                            seq length: 8
seq length: 14
 protein search, using sw model
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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                                                                                                                                                                                                                                                                           SUMMARIES
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                                               Rhesus blood group
T-cell receptor be
dodecenoyl-CoA Del
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| S32471 | C30503 | S23926 | PC2372 | A24867 | S78420 | PT0324 | B27867 | S37141 | PC7079 | PT0252 | A01250 | PS0443 | E60396 | S63492 | A44818 |
| lymnaDFamide 1 - g | Ig gamma-2b chain | major glycoprotein | 58K heat shock pro | scyliorhinin I - s | ribosomal protein | Ig heavy chain CRD | homeotic protein U | rpsA protein - Erw | unidentified 27.2K | Ig heavy chain CRD | angiotensin precur | potassium channel | antigen 7H8/2 - ma | dissimilatory sulf | extracellular lipa |

ALIGNMENTS

R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991 A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102 A;Accession: PT0254 R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991 A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102 A;Accession: PT0331 A;Molecule type: DNA A;Residues: 1-13 <YAM> A;Experimental source: B lymphocyte C;Keywords: heterotetramer; immunoglobulin Ig heavy chain CRD3 region (clone J2-121) - human (fragment) C;Species: Homo sapiens (man) C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996 C;Accession: PT0331 A;Molecule type: DNA A;Residues: 1-14 <YAM> A;Experimental source: B lymphocyte 밁 片 Ś C; Keywords: heterotetramer; Ig heavy chain CRD3 region (clone 2-115A) - human (fragment) C;Species: Homo sapiens (man) C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996 C;Accession: PT0254 RESULT 1 Matches Matches Query Match Best Local Similarity \vdash 2 YKNSKFHR 9 σ 4; Similarity 4; Conserv GYKNSKF 7 GYNNPPF 12 YDSSGYHR 14 Conservative Conservative 41.7%; 38.3%; 57.1%; immunoglobulin 0, 2; Mismatches Score 23; DB 2; Pred. No. 4e+02; Score 25; DB 2; Pred. No. 1.8e+02; Mismatches DB 2; Length 13; Length 14; Indels Indels 0; 0 Gaps Gaps 0 0

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RESULT PH1454

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C;Species: Mus musculus (house mouse)

C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 11-Apr-1995

C;Accession: PH1454

R;Casanova, J.L.; Martinon, F.; Gournier, H.; Barra, C.; Pannetier, C.; Regnault, A.; Kc

J. Exp. Med. 177, 811-820, 1993

A;Title: T cell receptor selection by and recognition of two class I major histocompatib

A;Reference number: PH1430; MUID:93171821; PMID:8436911

A;Accession: PH1454

A;Molecule type: mRNA

A;Residues: 1-12 <CAS>

A;Experimental source: cytolytic T-lymphocyte

C;Superfamily: immunoglobulin homology

C;Keywords: receptor; T-cell
                                                                                                                                                                                                                                                      R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L. J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility allelic exclusion and antigen-specific repertoire.
A;Reference number: PH0746; MUID:92078846; PMID:1836010
A;Accession: PH0787
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PH0787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           telomeric and tetraplex DNA binding protein qTBP42 VIII - rat (fragment) C;Species: Rattus norvegicus (Norway rat)
C;Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 07-May-1 C;Accession: PC4377
R;Sarig, G.; Weisman-Shomer, P.; Fry, M.
Biochem. Biophys. Res. Commun. 237, 617-623, 1997
A;Title: Telomeric and tetraplex DNA binding properties of qTBP42: A hom. A;Reference number: PC4371; MUID:97445086; PMID:9299414
A;Accession: PC4371
A;Accession: PC4377
                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 1-13 < CAS>
A; Cross-references: EMBL: X60891
A; Experimental source: T lymphocyte
C; Keywords: T-cell receptor
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PC4377
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C; Comment: '
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Best Local S
Matches 4
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;Species: Mus musculus (house mouse)
;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
;Accession: PH0787
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Pred. No. 5.7e
0; Mismatches
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Pred. No.
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Pred. No. 6.2e+02;
1; Mismatches 2
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R;Nakano, N.; Kikutani, H.; Nishimoto, H.; Kishimoto, T. J. Exp. Med. 173, 1091-1097, 1991
A;Title: T cell receptor V gene usage of islet beta cell-reactive T cells is A;Reference number: PT0209; MUID:91217621; PMID:1902501
A;Accession: PT0211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: protein
A; Residues: 1-10 < KAN>
R; Kozawa, H.; Hino, J.; Minamino, N.; Kangawa,
Biochem. Biophys. Res. Commun. 177, 588-595, 19
A; Title: Isolation of four novel tachykinins fi
A; Reference number: JE0426; MUID: 91254337; PMII
A; Accession: JE0427
                                                                                                                                                                                    T-cell receptor alpha chain V-J region (4-1-G.4) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 30-May-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      submitted to JIPID, February 1994
A,Description: Phylogenetic diversity
A,Reference number: JP0042
A,Accession: JP0073
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A;Title: Isolation of four novel tachykinins from frog (Rana catesbeiana) brain and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ranatachykinin B - bullfrog
C;Species: Rana catesbeiana (bullfrog)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993
C;Accession: B61033; JE0427
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A; Accession: B61033
                   A; Molecule type: mRNA
A; Residues: 1-11 < NAK>
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;Residues: 1-11 <NAK>
;Keywords: T-cell receptor
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                                                                                                                                                               Accession: PT0211
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100.0%; Pred. No. 2.
Live 0; Mismatches
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1992
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PMID:2043143
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cytochrome-c oxidase (EC 1.9.3.1) chain VIb, hepatic - rat (fragment) C;Species: Rattus norvegicus (Norway rat) C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 07-May-1999 C;Accession: S65381 R;Schaegger, H.; Noack, H.; Halangk, W.; Brandt, U.; von Jagow, G. Eur. J. Biochem. 230, 235-241, 1995 A;Title: Cytochrome-c oxidase in developing rat heart. Enzymic properties ar A;Reference number: S65372; MUID:95324529; PMID:7601105
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R; Volknandt, W.; Pevsner, J.; Elferink, L.A.; Scheller, R.H. FEBS Lett. 317, 53-56, 1993
A; Title: Association of three small GTP-binding proteins with cholinergic synaptic vesi A; Reference number: S29485; MUID: 93154521; PMID: 8428634
A; Accession: S29486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTP-binding protein o-rab3 - electric ray (Discopyge ommata) (fragment) C;Species: Discopyge ommata C;Species: Discopyge ommata C;Date: 22-Nov-1993 #sequence_revision 27-Feb-1997 #text_change 13-Mar-1997 C;Date: 32-Nov-1993 #sequence_revision 27-Feb-1997 #text_change 13-Mar-1997 C;Date: 329486
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A;Accession: S22236
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A;Title: Purification and characterization of two lipoxygenase isoenzymes from gangle A;Reference number: S21772; MUID:92207997; PMID:1554746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: S22236
R;Doderer, A.; Kokk
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C;Species: Hordeum vulgare (barley)
C;Date: 19-Mar-1997 #sequence_revision 24-Mar-1999 #text_change 24-Mar-1999
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;Residues: 1-14
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Pred. No. 1.3e+03;
1; Mismatches 0
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1; Mismatches
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1.6e+03;
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1.6e+03;
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   leucokinin VI - Madeira cockroach
C;Species: Leucophaea maderae (Madeira cockroach)
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990
C;Accession: JS0316
R;Holman, G.M.; Cook, B.J.; Nachman, R.J.
                                                                                                         RESULT
JS0316
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-13 <YAS>
C;Superfamily: unassigned
C;Keywords: skin
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C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 18-Aug-2000
C;Accession: A61361
R;Yasuhara, T.; Hira, M.; Nakajima, T.; Yanaihara, N.; Yanaihara, C.; Hashimoto, T.; S
Chem. Pharm. Bull. 21, 1388-1391, 1973
A;Title: Active peptides on smooth muscle in the skin of Bombina orientalis Boulenger
A;Reference number: A61361; MUID:73256822; PMID:4732297
A;Accession: A61361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-8 <SCH>
C;Keywords: oxidoreductase
                                                                                                                                                                                                                                              C;Accession: S47358
R;Lehner, P.J.
submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recognition
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S47358
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                                                                                                                                                                                                                                                                                                                                   T-cell antigen receptor VJ junction beta chain - human C; Species: Homo sapiens (man)
                                                                                                                              A;Cross-references: EMBL:Z35682; NID:g527453; PIDN:CAA84751.1; C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A61361
                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-13 < LEH >
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                                                                                                                                                                                                                                                                                                      ;Species: Homo sapiens (man)
;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 05-Nov-1999
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Pred. No.
                                                               Score 19; DB 2;
Pred. No. 2.3e+03;
1; Mismatches 3
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Pred. No.
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#text_change 20-Jun-2000

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C; Species: Rana
C; Date: 24-May
C; Accession: As
R; BANERJEE, H.
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A;Title: Isolation, primary structure, and synthesis of leucokinins V and A;Reference number: JS0315
A;Accession: JS0316
A;Molecule type: protein
A;Residues: 1-8 <HOL>
C;Comment: Leucokinins, a family of cephalomyotropic peptides, stimulate c;CKeywords: amidated carboxyl end; cephalomyotropic peptide; pyroglutamic F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;8/Modified site: amidated carboxyl end (Gly) #status experimental
D58503
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PT0274

Ig heavy chain CRD3 region (clone 3-109B) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-
C;Date: 70-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-
C;Accession: PT0274
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.
J. Exp. Med. 173, 395-407, 1991
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy
A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0274
A;Molecule type: DNA
A;Molecule type: DNA
A;Molecule type: DNA
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A; Residues: 1-12 < YAM>
A; Experimental source: B lymphocyte
C; Keywords: heterotetramer; immunoglobulin
                  RESULT 17
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A; Accession: A59387
                                                                                                                                                                                                                                                                                                                                                       submitted to the Protein Sequence A; Description: Identification and
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A59387
VCAM-1 5'UTR binding protein - Rana pipiens (fragment)
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A; Residues: 1-13 <BAN>
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Best Local S
Matches 3
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;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change
;Accession: A59387
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Best Local S
Matches 3
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VCAM-1 5' untranslated region binding protein with a probable translation inhibi
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1; Mismatches
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Pred. No.
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Pred. No. 2.8e+05;
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                                                                                                                                                  Mismatches
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. 3.6e+03;
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3.3e+03;
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cytochrome-c oxidase (EC 1.9.3.1) chain VII b, ca C;Species: Rattus norvegicus (Norway rat) C;Date: 12-Feb-1998 #sequence revision 20-Feb-199 C;Accession: S65387; S65386 R;Schaegger, H.; Noack, H.; Halangk, W.; Brandt, Eur. J. Biochem. 230, 235-241, 1995 A;Title: Cytochrome-c oxidase in developing rat h, R;Reference number: S65372; MUID: 95324529; PMID: 7, Section: S65387
                                                                                                                                      telomeric and tetraplex DNA binding protein qTBP42 IV - C;Species: Rattus norvegicus (Norway rat)
C;Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text
C;Accession: PC4374
R;Sarig, G.; Weisman-Shomer, P.; Fry, M.
Biochem. Biophys. Res. Commun. 237, 617-623, 1997
A;Title: Telomeric and tetraplex DNA binding properties
A;Reference number: PC4371; MUID:97445086; PMID:9299414
                                                                                                                                                                                                                                                                                    RESULT 19
PC4374
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R;Binette, J.P.; Binette, M.B.
submitted to the Protein Sequence Database,
submitted to the proteins of kidney and g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         translation elongation factor EF-Tu - unidentified bacterium C;Species: unidentified bacterium C;Species: unidentified bacterium C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_chan
                                                                                       A; Molecule type: protein A; Residues: 1-10 <SAR>
                                                                                                                       A; Accession: PC4374
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                 28.3%;
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Score 17; DB Pred. No. 4.3e 2; Mismatches
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                                                                       telomeric DNA
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S36678

Godecenoyl-CoA Delta-isomerase (EC 5.3.3.8) / enoyl-CoA hydratase (EC 4.2.1.17) / 3-h; C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 07-Nov-1997 #text_change 24-Jul-1998
C;Accession: S36678
R;Chen, N.; Crane, D.I.
Biochem. J. 283, 605-610, 1992
A;Title: Induction of the major integral membrane protein of mouse liver peroxisomes A;Reference number: S21285; MUID:92246895; PMID:1575703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cell receptor beta chain V-D-J region (clone 10) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: pH0939
R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
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A; Residues: 1-11 <GOL>
A; Experimental source:
C; Keywords: T-cell rece
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A;Title: Analysis of T cell receptor beta chains in Lewis
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A;Title: Organization of the gene (RHCE) encoding the human blood group RhCcEe antigens A;Reference number: I54193; MUID:94245182; PMID:8188244
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                  A; Molecule type: protein A; Residues: 1-14 < CHE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 1-11 <RES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Experimental source: complete Freund's adjuvant-immunized lymph Keywords: T-cell receptor
Experimental source: liver
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Pred. No. 4.7e+03;
1; Mismatches 1
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Pred. No. 4.7e+03;
3; Mismatches
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RESULT
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R;Lorenz, M.W.; Kellner, k.; Accession
J. Biol. Chem. 270, 21103-21108, 1995
J. Biol. Chem. 270, 21103-21108, 1995
A;Title: A family of neuropeptides that inhibit juvenile
N;Alternate names: locustatachykinin IV
C;Species: Locusta migratoria (migratory locust)
C;Date: 31-Mar-1993 #sequence_revision 31-Mar-1993
C;Accession: B60073
R;Schoofs, L.; Holman, G.M.; Hayes, T.K.; Kochansky
                                                                                                                                                                                                                                                                                                                                                                                   A;Experimental source: pituitary A;Note: sequence extracted from C;Keywords: oxidoreductase
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A;Pathway: fatty acid beta-oxidation
C;Keywords: carbon-oxygen lyase; fatt
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                                                                                                      tachykinin IV - migratory locust
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2; Conserv
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Pred. No. 6.1e+03
1; Mismatches
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Pred. No. 2.8e+05;
3; Mismatches 0
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L.; Holman, G.M.; Hayes, T.K.; Kochansky,

J.P.; Nachman,

R.J.; De

Loof,

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\epsilontranslation
                                RESULT 28
S01222
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S23373
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A; Title: Locustatachykinin III and IV: two additional insect neuropeptides with homology A; Reference number: A60073; MUID:91219696; PMID:2132575
A; Accession: B60073
A; Molecule type: protein
A; Residues: 1-10 <SCH>
C; Superfamily: tachykinin
C; Keywords: amidated carboxyl end; neuropeptide; tachykinin
C; Keywords: amidated carboxyl end (Arg) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0243
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A; Residues: 1-11 < PLU>
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                                                                                                                                                                                                                                                                                                                                                                                                           T-cell receptor alpha chain J region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C;Accession: S23373
C;Accession: S23373
R;Pluschke, G.; Ricken, G.; Taube, H.; Kroninger, S.; Melchers, I.; Peter, H.H.; Eichman Eur. J. Immunol. 21, 2749-2754, 1991
A;Title: Biased T cell receptor V(alpha) region repertoire in the synovial fluid of rheu A;Reference number: S23364; MUID:92037820; PMID:1657615
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A;Residues: 1-10 <YAM>
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0243
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Best Local S
Matches 3
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Best Local S
Matches 3
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;Keywords: T-cell receptor
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Paramonds: heterotetramer; immunoglobulin
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Best Local Similarity
Matches 3; Conserv
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        elongation factor EF-Tu
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3; Conserve
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Pred. No. 6.6e+03;
1; Mismatches 1
                                                                                                                                                                                                Score 16; DB 2;
Pred. No. 7.3e+03;
2; Mismatches 2
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        Pseudomonas aeruginosa (fragment)
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c;Species: Xenopus laevis (African clawed frog)
C;Species: Xenopus laevis (African clawed frog)
C;Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
C;Accession: S10624
R;Wallace, R.A.; Hoch, K.L.: Carnaval:
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A44818
                                                                                                                                                                                                          R;Gilbert, E.J.; Cornish, A.; Jones, C.W.
J. Gen. Microbiol. 137, 2223-2229, 1991
A;Title: Purification and properties of extracellular lipase from A;Reference number: A44818; MUID:92085040; PMID:1748875
A;Accession: A44818
A;Status: preliminary; not compared with conceptual translation A;Molecule type: nucleic acid
A;Residues: 1-13 <GIL>
                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Pseudomonas aeruginosa
C;Date: 31-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 31-Oct-1997
C;Accession: A44818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Wallace, R.A.; Hoch, K.L.; Carnevali, O. J. Mol. Biol. 213, 407-409, 1990
A;Title: Placement of small lipovitellin subunits within the vitellogenin precursor in A;Reference number: S10624; MUID:90278951; PMID:2352275
A;Accession: S10624
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C;Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 26-Aug-1999
C;Accession: S01222
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C;Superfamily: translation elongation factor Tu; translation elongation factor
C;Keywords: GTP binding; protein biosynthesis
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Nucleic Acids Res. 16, 7193, 1988

A;Title: A fragment of the Pseudomonas aeruginosa genome contains five tRNA genes,
A;Reference number: S01222; MUID:88303352; PMID:3136442
                                                                                                                                           A;Note: sequence extracted from Note Dacker C;Superfamily: Pseudomonas triacylglycerol
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A; Residues: 1-12 < HUG>
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Best Local S
Matches 3
                                                                                                                                                                   Experimental source: strain TE3285
Note: sequence extracted from NCBI backbone (NCBIP:70395)
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Best Local Similarity
Matches 3; Conserv
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3; Conserv
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larity 30.0%;
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75.0%;
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Pred. No. 8.7e+03;
3; Mismatches 4
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Pred. No. 8e+03;
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Pred. No. 8e+03;
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antigen 7H8/2 - malaria parasite (Plasmodium falciparum) (fragments)
C;Species: Plasmodium falciparum
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 09-Jun-2000
C;Accession: E60396
R;Limpaiboon, T.; Taylor, D.W.; Jones, G.; Geysen, H.M.; Saul, A.
Southeast Asian J. Trop. Med. Public Health 21, 388-396, 1990
A;Title: Characterization of a Plasmodium falciparum epitope recognized by a
                                                                                                                                                                                                                                                                                                                             R; Adelman, J.P.; Shen, K.Z.
Neuron 9, 209-216, 1992
A; Title: Calcium-activated :
A; Reference number: JH0697;
A; Accession: PS0443
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$63492
dissimilatory sulfite reductase beta chain, soluble - Desulfovibrio desulfuricans
C; Species: Desulfovibrio desulfuricans
C; Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
                                                                                                                                                                                           A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 1-13 <ADE>
C;Comment: This potassium channel is activated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            potassium channel protein Slo G3 - fruit fly (Drosophila melanogaster) (fragment) C;Species: Drosophila melanogaster C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Feb-1997 C;Accession: PS0443
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A;Accession: E60396
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A;Molecule type: protein
A;Residues: 1-13 <STE>
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C;Accession: S63492
R;Steuber, J.; Arendsen, A.F.; Hagen, W.R.; Kroneck, P.M.H.
R;Steuber, J.; Arendsen, A.F.; Hagen, W.R.; Kroneck, P.M.H.
Bur. J. Biochem. 233, 873-879, 1995
Bur. J. Biochem. 233, 873-879, 1995
A;Title: Molecular properties of the dissimilatory sulfite reductase from Desulfovibrio A:Reference number: S63489; MUID:96085152; PMID:8521853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Cross-references: GB:M31305
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Best Local S
Matches 3
                                                            ;Cross-references: FlyBase:FBgn0003429;Keywords: alternative splicing; ion c
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;Residues: 1-13 <LIM>
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26.7%;
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  26.7%;
                                                                                                                                                                                                                                                                                                                                                                potassium channels expressed from cloned complementary DNAs.
; MUID:92360298; PMID:1497890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16; DB:
Pred. No. 8.7e.
3; Mismatches
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Pred. No. 8.7e
0; Mismatches
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  Score
                                                               ion channel; potassium channel;
16;
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8.7e+03;
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8.7e+03;
DB
'n
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Length 13
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                                                                  transmembrane
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                                                               protein
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                                                                                                                                                                                                                                                                                                                                                                                                 RESULT
PC7079
                              A;Title: Proteome analysis of mouse brain: Two-dimensional electrophoresis profiles A;Reference number: PC7072
A;Accession: PC7079
A;Molecule type: protein
A;Residues: 1-14 <TSU>
                                                                                                                                                                                     unidentified 27.2K protein - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 18-Aug-2000
C;Accession: PC7079
R;Tsugita, A.; Kawakami, T.; Uchida, T.; Sakai, T.; Kamo, M.; Matsui, T.; Wallstrophoresis 21, 1853-1871, 2000
Electrophoresis 21, 1853-1871, 2000
Almittophoresis 21, 2000
Almitto
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A;Experimental source:
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strain

C57BL/6Cr Slc,

male; brain,

striatum

Watanabe,

of f ۲.,

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C;Accession: A92775; A01250
R;Skeggs Jr., L.T.; Kahn, J.R.; Lentz, K.; Shumway, N.P.
J. Exp. Med. 106, 439-453, 1957
A;Reference number: A92775
A;Accession: A92775
A;Accession: A92775
A;Molecule type: protein
A;Residues: 1-14 <SKE>
C;Superfamily: Serpin
C;Keywords: blood pressure control; hormone; vasoconstrictor
F;1-10/Product: angiotensin I #status experimental <AN1>
F;1-8/Product: angiotensin II #status experimental <AN2>
                                                                                                                                                                                                                                                                                                      Ig heavy chain CRD3 region (clone 2-109D) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0252
                                                                                                                                          A;Molecule type: DNA
A;Residues: 1-14 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunogl
                                                                                                                                                                                                                R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0252
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Best Local :
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Best Local Similarity
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Matches 3; Conserv
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                                                                   Similarity
3; Conserv
                                 YKNSKFHRV 10
YDSSGYYYV
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                                                                                       33.3%;
                                                                                                         26.7%;
                                                                                                                                        immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 16; DB Pred. No. 9.4e
2; Mismatches
                                                                   Score 16; DB Pred. No. 9.4e 3; Mismatches
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در
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2; Mismatches 2;
                                                                                        9.4e+03
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                                                                                                         DB 2;
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                                                                                                      Length 14;
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C; Keywords: brain

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R;Saari, G.; Bienz, M.
EMBO J. 6, 1775-1779, 1987
A;Title: The structure of th
A;Reference number: A91072
A;Accession: B27867
A;Molecule type: mRNA
A;Residues: 1-8 <SAA>
C;Genetics:
Ig heavy chain CRD3 region (clone J2-106C) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-C;Accession: PT0324
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy
A;Reference number: PT0222; MUID:91108337; PMID:1899102
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S37141
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                                                                                                                                                                                                                                                                                                                                             A;Gene: FlyBase:Ubx
A;Cross-references: FlyBase:FBgn0003944
C;Keywords: DNA binding; nucleus; trans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 38
B27867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rpsA protein - Erwinia chrysanthemi
C;Species: Erwinia chrysanthemi
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-199
C;Accession: S37141
R;Douillie, A.; Toussaint, A.; Faelen, M.
submitted to the EMBL Data Library, August 1993
A;Description: Identification of the integration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     homeotic protein Ultrabithorax - fruit fly (Drosophila melanogaster) (fragment) C;Species: Drosophila melanogaster C;Species: Drosophila melanogaster C;Date: 08-Mar-1989 #sequence_revision 08-Mar-1989 #text_change 23-Feb-1997 C;Accession: B27867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Molecule type: DNA
A;Residues: 1-8 <DOU>
A;Cross-references: EMBL:X74750; NID:g399669; PIDN:CAA52769.1; PID:g581108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Description: Identificat:
A; Reference number: S37139
A; Accession: S37141
A; Status: preliminary
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Best Local S
Matches 3
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3; Conserv
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50.0%;
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Pred. No. 2.8e
1; Mismatches
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Pred. No. 9.4e+03;
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Pred. No. 2.
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                                                                                       #text_change 16-Aug-1996
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                                                        Caton, A.J.; Rovera, G.
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RESULT
S78420
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A;Molecule type: DNA
A;Residues: 1-9 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                           R; Waugh, D.; Wang, Y.; Hazon, N.; Balment, Eur. J. Biochem. 214, 469-474, 1993
A; Title: Primary structures and biological A; Reference number: S33300; MUID:93292508; A; Accession: S33301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT
A24867
                                                                                                                                      C;Superfamily: unassigned animal peptides C;Keywords: amidated carboxyl end; neuropeptide F;10/Modified site: amidated carboxyl end (Met)
                                                                                                                                                                                                                                                                                                                            A;Accession: A24867
A;Molecule type: protein
A;Residues: 1-10 <CON>
                                                                                                                                                                                                                                                                                                                                                                                       R;Conlon, J.M.; Deacon, C.F.; O'Toole, L.; Thim, L. FEBS Lett. 200, 111-116, 1986
A;Title: Scyliorhinin I and II: two novel tachykinins from A;Reference number: A91359; MUID:86192829; PMID:2422058
                                                                                                                                                                                                                                                                                                                                                                                                                                                          scyliorhinin I - smaller spotted catshark C;Species: Scyliorhinus canicula (smaller spotted catshark, smaller spotted dogfish) C;Date: 02-Jun-1988 #sequence_revision 02-Jun-1988 #text_change 18-Aug-2000 C;Accession: A24867; S33301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: protein
A; Residues: 1-9 < GOL>
A; Note: the protein is designated as mitochondrial ribosomal protein
C; Keywords: mitochondrion; protein biosynthesis; ribosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the Protein S A; Reference number: S78411 A; Accession: S78420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Accession: S78420 R; Goldschmidt-Reisin, S.; Graack,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ribosomal protein RL41, mitochondrial [validated] - rat (fragment) C;Species: Rattus norvegicus (Norway rat) C;Date: 25-Feb-1998 #sequence_revision 13-Mar-1998 #text_change 21
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                                                  Query Match
Best Local Similarity
Thes 3; Conserve
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A;Residues: 1-10 <WAU>
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                                                                                                                                                                                                                            A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                      A; Reference number: A91359;
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                                  KNSKFH 8
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50.0%;
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33.3%;
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60.0%;
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                                                                    Score 15; DB 2;
Pred. No. le+04;
1; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                   R.J.; Conlon,
                                                                                                                                                                                                                                                                  activities of PMID:7685693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     February 1998
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                                                                                                                                         #status experimental
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Ig gamma-2b chain C region (F5.5.1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 16-Aug-1996
C;Accession: C30503
R;Gilmore, G.L.; Bard, J.A.; Birshtein, B.K.
J. Immunol. 141, 1754-1761, 1988
A;Title: DNA rearrangements affecting both variable and constant regions of A;Reference number: A30503; MUID:88315788; PMID:2842402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 43
S23926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Kim, D.H.; Kanno, C.; Mizokami, Y.
Biochim. Biophys. Acta 1122, 203-211, 1992
A;Title: Purification and characterization of major glycoproteins, A;Reference number: S23926; MUID:92353107; PMID:1643094
A;Accession: S23926
A;Molecule type: protein
A;Residues: 1-11 <KIM>
C;Keywords: glycoprotein; milk; blocked amino end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Matsuno, K.; Miyamoto, T.; Yamaguchi, K.; Sayed, M.A.; Kajiwara, T.; Hatano, S. Biosci. Biotechnol. Biochem. 59, 231-235, 1995
A;Title: Identification of DNA-binding proteins changed after induction of sporulation A;Reference number: PC2369; MUID:95218265; PMID:7766022
A;Accession: PC2372
                                 Ś
                                                                                                                                                                                               A;Reference number: A30503;
A;Accession: C30503
A;Molecule type: mRNA
A;Residues: 1-12 <GIL>
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C;Species: Bos primigenius taurus (cattle)
C;Species: 19-Mar-1997 #sequence_revision 01-Feb-1999 #text_change 01-Feb-1999
C;Date: 19-Mar-1997 #sequence_revision 01-Feb-1999 #text_change 01-Feb-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 42
PC2372
B
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A;Residues: 1-11 <MAS>
C;Keywords: heat shock; molecular chaperone; stress-induced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Status: preliminary
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Best Local Similarity 60.0
                                                                                                                                                       Experimental source: myeloma cell line Keywords: immunoglobulin
                                                                                               Query Match
Best Local (
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les 3; Conserv
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Pred. No. 1.
                                                                                               Score 15; DB 2;
Pred. No. 1.2e+04;
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Pred. No. 1.1e+04
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1.1e+04;
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                                                                                                                 Length 12;
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R; Johnsen, A.H.; Rehfeld, J.F.
Eur. J. Biochem. 213, 875-879, 1993
A; Title: LymnaDFamides, a new family of neuropeptides from A; Reference number: S32471; MUID:93238777; PMID:8477756
A; Accession: S32471
                                                                                                                                               A;Cross-references: PIDN:AAB26362.1; PID:g299829
A;Experimental source: ganglia
C;Keywords: amidated carboxyl end; neuropeptide
F;13/Modified site: amidated carboxyl end (Phe) #status predicted
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S32471
                               Ś
                                                                                                                                                                                                                                                                                                                                                                               C;Species: Lymnaea stagnalis (great pond snail)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
                                                                                                                                                                                                                           A;Molecule type: protein
A;Residues: 1-13 <JOH>
                                                                                                                                                                                                                                                                                                                                                                                                                          lymnaDFamide 1 -
                                                                                        Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                  Accession: S32471;
                                                                          Matches
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                                                                       Similarity
3; Conserv
                                 NSKF 7
NSAF 10
                                                                                                                                                                                                                                                                                                                                                                                                                        great pond snail
                                                                          Conservative
                                                                                          25.0%;
75.0%;
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                                                                         Score 15; DB 2;
Pred. No. 1.3e+04;
0; Mismatches 1
                                                                                                          Length 13;
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Search completed: August 30, 2004, 10:58:49 Job time: 4.30743 secs

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Title:
Perfect score:
Sequence:
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No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                               Score
      seq length: 8
seq length: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                   Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*

12: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*

13: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*

14: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep:*

15: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

16: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

17: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*

16: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

17: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

18: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
                                                                                                                                                                                                                                                              Query
Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             August 30, 2004, 10:49:45; Search time 13.4527 Seconds (without alignments)
257.253 Million cell updates/sec
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Gapop 10.0 , Gapext 0.
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     GYKNSKFHRVI 11
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                                                                                                                                                                                                                                                               Length
      GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd
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 6 US-10-447-161-84
6 US-10-788-016-1
14 US-10-105-232-266
14 US-10-189-437-253
US-09-226-666-1
13 US-09-226-666-13
14 US-09-226-666-13
15 US-09-226-666-16
16 US-10-601-953-588
17 US-10-601-953-588
18 US-10-322-266-460
19 US-09-780-053-158
19 US-09-780-053-304
19 US-09-780-053-304
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Sequence 84, Appli
Sequence 1, Appli
Sequence 266, App
Sequence 253, App
Sequence 1, Appli
Sequence 13, Appl
Sequence 16, Appl
Sequence 16, App
Sequence 588, App
Sequence 460, App
Sequence 158, App
Sequence 158, App
Sequence 158, App
Sequence 112, Appl
Sequence 180, App
Sequence 304, App
Sequence 399, App
                                                                                                                                                                                                                                                              Description
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ALIGNMENTS

RESULT 1 US-10-447-161-84

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RESULT 2
US-10-788-016-1
; Sequence 1, Application US/10788016
; Publication No. US20040141992A1
; GENERAL INFORMATION:
; APPLICANT: ITOH, Kyogo
; TITLE OF INVENTION: Desensitizers
                                                                                                                                                                                   밁
                                                                                                                                                                                                                                                                                                                               ; FEATURE:
; OTHER INFORMATION: Synthetic Peptide
US-10-447-161-84
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                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Wang, Rong-fu
TITLE OF INVENTION: Mutant Fibronectin and Tumor Metastasis
FILE REFERENCE: HO-P02484US1
CURRENT APPLICATION NUMBER: US/10/447,161
CURRENT FILING DATE: 2003-05-28
PRIOR APPLICATION NUMBER: 60/383,530
PRIOR FILING DATE: 2002-05-28
NUMBER OF SEQ ID NOS: 148
SOFTWARE: PatentIn version 3.1
SEQ ID NO 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 84, Application US/10447161
Publication No. US20040023314A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                       Local Similarity hes 6; Conserv
                                                                                                                                                                              6 KFHRVI 11
||||||
1 KFHRVI 6
                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                      53.3%; Score 32; DB 16; 100.0%; Pred. No. 1.2e+06
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; Sequence 253, Application US/10189437
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                                                                                                                                             8
                                                                                                                                                                                                                                                                         ; FEATURE:
; NAME/KEY: MOD_RES
; LOCATION: (8)
; OTHER INFORMATION: gly or gln
US-10-105-232-266
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                    RESULT 4
US-10-189-437-253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/105,232
CURRENT FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR APPLICATION NUMBER: 08/198,139
PRIOR FILING DATE: 1994-02-17
PRIOR FILING DATE: 1994-02-17
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Best Local S
Matches 4
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SEQ ID NO 266
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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CURRENT APPLICATION NUMBER: US/10/788,016
CURRENT FILING DATE: 2004-02-26
PRIOR APPLICATION NUMBER: PCT/JP02/08641
PRIOR FILING DATE: 2002-08-28
PRIOR APPLICATION NUMBER: JP P2001-260046
PRIOR FILING DATE: 2001-08-29
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BOGOCH, SAMUEL APPLICANT: BOGOCH, ELENO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKIN PEPTIDES IN RAPID REPLICATION OF GLIOMA CELLS
TITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS
FILE REFERENCE: 09425-46904
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Influenza virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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Similarity 100.0%; Pred. No. 1.2e+06;
6; Conservative 0; Mismatches 0;
                                                                                                                                                 NSKFHRV 10
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o. US20030180328A1
                                                                                                                                                                                              Conservative
                                                                                                                                                                                                              45.0%;
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Pred. No. 3.6e+02;
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APPLICANT: Spacciapoli, Peter
APPLICANT: Spacciapoli, David M.
APPLICANT: Friden, Phillip M.
FITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
CURRENT APPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/189,437
CURRENT FILING DATE: 2002-07-08
PRIOR APPLICATION NUMBER: 10/105,232
PRIOR FILING DATE: 2002-03-26
PRIOR PILING DATE: 2002-03-26
PRIOR FILING DATE: 2001-10-26
PRIOR PILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 729
SOFTWARE: PATENTIN 2.1
SEQ ID NO 253
                                     RESULT 6
US-09-226-666-13
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US-09-226-666-1
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Best Local Similarity
Grandes 6; Conserva
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                                                                                                                                                                                                                                                                       OTHER INFORMATION: Internal fragment US-09-226-666-1
Sequence 13, Application US/09226666A Patent No. US20020077282A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/09226666A Patent No. US20020077282A1 GENERAL INFORMATION:
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Best Local Similarity 57.1%;
Matches 4; Conservative
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APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKIN PEPTIDES AND ANTIBODIES THEREFORE
FILE REFERENCE: 09425/46905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 11
TYPE: PRT
ORGANISM: Influenza virus
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NAME/KEY: MOD_RES
LOCATION: (8)
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                                                                                                                   6 GYKR-KFH 12
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Pred. No. 3.6e+02;
1; Mismatches 2
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GENERAL INFORMATION:

Spacciapoli,

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APPLICANT: Spacciapoli, Peter
APPLICANT: Rothstein, David M.
APPLICANT: Friden, Phillip M.
TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
CURRENT APPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16
LENGTH: 12
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                                                                                                                                                                                                                                                           RESULT 8
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Best Local S
Matches 6
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TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
CURRENT APPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bequence 16, Application US/09226666A Patent No. US20020077282A1 GENERAL INFORMATION:
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Best Local :
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APPLICANT: Quay, Steven
APPLICANT: El Shafy, Mohammed Abd
APPLICANT: Gupta, Malini
APPLICANT: de Meireles, Jorge
TITLE OF INVENTION: Compositions and Methods for Enhanced
TITLE OF INVENTION: Mucosal-Delivery of Interferon Beta
FILE REFERENCE: 02-02US
CURRENT APPLICATION NUMBER: US/10/462,452
CURRENT FILING DATE: 2003-06-16
                                                                                                                                                                                                    Sequence 459, Application US/10462452 Publication No. US20040037809A1
                                                                                                                                                                                  GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Derived from Homo sapiens 09-226-666-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Derived from Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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ORGANISM: Artificial Sequence
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75.0%;
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Pred. No. 4.8e+02;
0; Mismatches 1;
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APPLICANT: Quay, Steven C.

APPLICANT: Quay, Steven C.

TITLE OF INVENTION: Compositions And Methods For Modulating Physical Composition of Invention: Junctional Adhesion Molecules For Enhanced ITITLE OF INVENTION: Therapeutic Compounds

FILE REFERENCE: 02-03US

CURRENT APPLICATION NUMBER: US/10/601,953

CURRENT FILING DATE: 2003-06-24

PRIOR APPLICATION NUMBER: 60/392,512

PRIOR APPLICATION NUMBER: 60/392,512

PRIOR FILING DATE: 2002-06-28

NUMBER OF SEQ ID NOS: 900

SOFTWARE: Patentin version 3.2

SEQ ID NO 588
                                                                                                                                                                   RESULT 10
US-10-322-266-460
; Sequence 460, Application US/10322266
; Publication No. US20040115135A1
; GENERAL INFORMATION:
; GENERAL INFORMATION:
; TITLE OF INVENTION: Compositions And Methods For Enhanced Mucosal Delivery Of Peptide
; TITLE OF INVENTION: YY And Methods For Treating And Preventing Obesity
; FILE REFERENCE: NPCIO567
; CURRENT APPLICATION NUMBER: US/10/322,266
; CURRENT FILING DATE: 2002-12-17
; NUMBER OF SEQ ID NOS: 797
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 460
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                                                         ; OTHER INFORMATION: Synthetic construct US-10-322-266-460
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 459
LENGTH: 8
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Query Match
Best Local Similarity
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                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Synthetic construct
                                                                                                                                                        ENGTH:
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o. US20040077540A1
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Pred. No.
 Score 26;
Pred. No.
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Pred. No.
 DB 16;
1.2e+06;
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                   Length
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Matches

4; Conservative

Mismatches

7:

Indels

0

Gaps

0

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APPLICANT: Elana Levin
APPLICANT: Steve Chappell Mitchell
APPLICANT: Aya Jakobovits
TITLE OF INVENTION: 83P5G4: A TISSUE SPECIFIC PROTBIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCER
FILE REFERENCE: 129.5U1
CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 158
                                                                                                                                                                                                                       APPLICANT: Universit, t Zurich
FILE OF INVENTION: Calcium binding proteins
FILE REFERENCE: Calsyntenins
CURRENT APPLICATION NUMBER: US/10/380,705
CURRENT FILING DATE: 2003-03-14
PRIOR APPLICATION NUMBER: EP 00810830.0
PRIOR FILING DATE: 2000-09-14
NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 12
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                                                                                                                                                       US-10-380-705-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Homo Sapiens US-09-780-053-158
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US-09-780-053-158
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Best Local S
Matches 4
                                                                       Query Match 43.:
Best Local Similarity 57.:
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 12, Application US/10380705
Publication No. US20040019919A1
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APPLICANT: Daniel E.H. Afar
APPLICANT: Pia M. Challita-Eid
                                                                                                                                                                      TYPE: PRT
ORGANISM: Gallus gallus
                                                                                                                                                                                                              LENGTH: 9
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Local Similarity 50.0%;
les 4; Conservative
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 YKPAEFH 8
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                                    YKNSKFH 8
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                                                                                           43.3%;
                                                                       Score 26; DB 15; Length 9; Pred. No. 1.2e+06; 2; Mismatches 1; Indels
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Pred. No. 1.2e+06;
3; Mismatches 1;
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RESULT 13

RESULT 15 US-09-780-053-399

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APPLICANT: Elana Levin
APPLICANT: Steve Chappell Mitchell
APPLICANT: Steve Chappell Mitchell
APPLICANT: Aya Jakobovits
IITLE OF INVENTION: 83P5G4: A TISSUE SPECIFIC PROTEIN
IITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCER
FILE REFERENCE: 129.5USU1
CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 304
LENGTH: 10
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; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 180
; LENGTH: 10
TYPE: PRT
; ORGANISM: Homo Sapiens
US-09-780-053-180
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US-09-780-053-304
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity 50.0%;
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rene S. Hubert
APPLICANT: Daniel E.H. Afar
APPLICANT: Pia M. Challita-Eid
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APPLICANT: Aya Jakobovits
TITLE OF INVENTION: 83P5G4: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCER
FILE REFERENCE: 129.5USU1
CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
CURRENT FILING DATE: 2001-02-09
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APPLICANT: Daniel E.H. Afar
APPLICANT: Pia M. Challita-Eid
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PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
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                                      1 GYKNSKFH 8
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GHONSTFY 8
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Steve Chappell Mitchell
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20020102640A1
                                                                               Conservative
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                                                                                              43,3%;
                                                                            Score 26; DB 9; L
Pred. No. 4.9e+02;
3; Mismatches 1;
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Pred. No. 4.9e+02;
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399, Application US/09780053 o. US20020102640A1

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RESULT 16
US-09-572
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CURRENT FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 1144
SOFTWARE: ProtPatent version 1.0
SEQ ID NO 759
LENGTH: 10
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 399
                                                                                                       Sequence 413, Application US/10462452 Publication No. US20040037809A1 GENERAL INFORMATION:
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Matches
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Best Local
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ORGANISM: Homo Sapiens
-09-780-053-399
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APPLICANT: Quay, Steven
APPLICANT: El Shafy, Mohammed Abd
APPLICANT: Gupta, Malini
APPLICANT: de Meireles, Jorge
TITLE OF INVENTION: Compositions and Methods for Enhanced
TITLE OF INVENTION: Mucosal-Delivery of Interferon Beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Proteom Ltd
TITLE OF INVENTION: Inter- complementary peptide listing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
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TITLE OF INVENTION: 83P5G4: A TISSUE SPECIFIC PROTEIN
TITLE OF INVENTION: HIGHLY EXPRESSED IN PROSTATE CANCE
FILE REFERENCE: 129.5USU1
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                                                                                                                                                                                                                                                                                                                   Local
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Steve Chappell Mitchell
Aya Jakobovits
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Daniel E.H. Afar
Pia M. Challita-Eid
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Pred. No. 4.9e
3; Mismatches
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Pred. No. 4.9e
1; Mismatches
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4.9e+02;
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Sequence 541, Application US/10601953

Publication No. US20040077540A1

GENERAL INFORMATION:
APPLICANT: Quay, Steven C.
ITILE OF INVENTION: Compositions And Methods For Modulating Physiology Of Epit TITLE OF INVENTION: Junctional Adhesion Molecules For Enhanced Mucosal Delive TITLE OF INVENTION: Therapeutic Compounds

FILE REFERENCE: 02-03US

CURRENT APPLICATION NUMBER: US/10/601,953

CURRENT FILING DATE: 2003-06-24

PRIOR APPLICATION NUMBER: 60/392,512

PRIOR APPLICATION NUMBER: 60/392,512

PRIOR FILING DATE: 2002-06-28

NUMBER OF SEQ ID NOS: 900

SOFTWARE: PatentIn version 3.2

SEQ ID NO 541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; FILE REFERENCE: 02-02US
; CURRENT APPLICATION NUMBER: US/10/462,452
; CURRENT FILING DATE: 2003-06-16
; PRIOR APPLICATION NUMBER: 60/393,066
; PRIOR FILING DATE: 2002-06-28
; NUMBER OF SEQ ID NOS: 790
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 413
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-462-452-413
           Sequence 414, Application US/10322266

Publication No. US20040115135A1

GENERAL INFORMATION:
APPLICANT: Quay, Steven C.
TITLE OF INVENTION: Compositions And Methods For EnlyTITLE OF INVENTION: YY And Methods For Treating And FILE REFERENCE: NPCI0567

CURRENT APPLICATION NUMBER: US/10/322,266

CURRENT FILING DATE: 2002-12-17

NUMBER OF SEQ ID NOS: 797

SOFTWARE: PatentIn version 3.2

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                                                                                                                                                                                                                                                       RESULT 19
US-10-322-266-414
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Best Local S
Matches 4
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Best Local Similarity
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TYPE: PRT
ORGANISM: Artificial Sequence
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ENGTH:
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Pred. No. 4.9e
1; Mismatches
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1; Mismatches
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Delivery Of
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TYPE: PRT
ORGANISM: Artificial Sequence

FEATURE:

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; OTHER INFORMATION: Synthetic construct US-10-322-266-414
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US-10-239-313A-642
                                                    SOFTWARE: PatentIn version 3.1
SEQ ID NO 116
LENGTH: 8
                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Cell Signaling Technology, Inc
APPLICANT: COMB, Michael J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                         APPLICANT: TAN, YI
TITLE OF INVENTION: PRODUCTION OF MOTIF-SPECIFIC AND CONTEXT-INDEPENDENT ANTIBODIES
FILE REFERENCE: CST-138 CIP2
CURRENT APPLICATION NUMBER: US/10/014,485A
CURRENT FILING DATE: 2002-03-18
PRIOR APPLICATION NUMBER: US 09/148,712
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: US 09/535,364
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/239,313A CURRENT FILING DATE: 2002-09-19 PRIOR APPLICATION NUMBER: FR 00/03711 PRIOR FILING DATE: 2000-03-23 PRIOR APPLICATION NUMBER: PCT 01/70772 PRIOR FILING DATE: 2001-03-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 697
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: MOLECULE OF PHARMACEUTICAL INTEREST COMPRISING AT ITS ITLE OF INVENTION: N-TERMINAL A GLUTAMIC ACID OR A GLUTAMINE IN THE FORM ITLE OF INVENTION: OF A PHYSIOLOGICALLY ACCEPTABLE STRONG ACID TLE REFERENCE: 343 727 - US
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ORGANISM: Homo sapiens FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                       Application US/10014485A
o. US20020168684A1
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100.0%; Pr
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Pred. No. 4.9e+02;
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APPLICANT: Bachmann, Martin
APPLICANT: Maurer, Patrick
APPLICANT: Spohn, Gunther
TITLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease
FILE REFERENCE: 1700.0330001
CURRENT APPLICATION NUMBER: US/10/289,456
CURRENT FILING DATE: 2002-11-07
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2002-07-19
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
COUNTABLE: Date: 170 Second 10 Second 10
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; LOCATION: (5)..(5)
; OTHER INFORMATION: PHOSPHORYLATION; serine at position 5 is
US-10-014-485A-116
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US-09-982-172-186
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US-10-289-456-91
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Best Local Similarity
Tatches 5; Conserve
                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/982,172
CURRENT FILING DATE: 2001-10-19
NUMBER OF SEQ ID NOS: 253
SOFTWARE: PatentIn version 3.1
SEQ ID NO 186
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Emil Israel Katz
TITLE OF INVENTION: PEPTIDES REPRESENTATIVE OF POLYPEPTIDES OF INTEREST AND ANTIBODIES
TITLE OF INVENTION: DIRECTED THEREAGAINST, AND METHODS, SYSTEMS AND KITS FOR GENERAT:
TITLE OF INVENTION: UTILIZING EACH
FILE REFERENCE: 01/22283
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                                                                                                                 LENGTH: 1
TYPE; PRT
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                                                                   ORGANISM: Artificial sequence
OTHER INFORMATION: Computer generated synthetic peptide
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Pred. No. 1.2e+06;
1; Mismatches 2
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Pred. No. 8.1e+02;
1; Mismatches 0
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; OTHER INFORMATION: Computer generated synthetic peptide US-09-982-172-245
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US-09-982-172-245
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Patent No. US20020137119A1
GENERAL INFORMATION:
APPLICANT: Emil Israel Katz
TITLE OF INVENTION: PEPTIDES REPRESENTATIVE OF POLYPEPTIDES OF INTEREST AND ANTIBODIE TITLE OF INVENTION: UTILIZING EACH
TITLE OF INVENTION: UTILIZING EACH
TITLE OF INVENTION: UTILIZING EACH
TITLE OF INVENTION UTILIZING EACH
TILE REFERENCE: 01/22283
CURRENT APPLICATION NUMBER: US/09/982,172
CURRENT APPLICATION NUMBER: US/09/982,172
CURRENT FILING DATE: 2001-10-19
NUMBER OF SEQ ID NOS: 253
SOFTWARE: Patentin version 3.1
SEQ ID NO 245
FEATURE:
NAME/KEY: MOD_RES
LOCATION: (8)...(8)
OTHER INFORMATION: 1
S-10-174-105A-153
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 09/148,712
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: US 09/535,364
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 193
                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: POSITIVE IDENTIFICATION OF PHOSPHO-PROTEINS USING MOTIF-SPECIFIC TITLE OF INVENTION: CONTEXT-INDEPENDENT ANTIBODIES COUPLED WITH DATABASE SEARCHING FILE REFERENCE; CST-138 CIP3
                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/174,105A CURRENT FILING DATE: 2002-06-18
                                                                                      FEATURE:
OTHER INFORMATION: Synthetic Peptide
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les 4; Conservative
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                  PHOSPHORYLATION;
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50.0%;
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Pred. No.
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Pred. No. 8.
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                    serine at
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APPLICANT: Spohn, Gunther
FITTLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease
FILE REFERENCE: 1700.0330001
CURRENT APPLICATION NUMBER: US/10/289,456
CURRENT FILING DATE: 2002-11-07
PRIOR APPLICATION NUMBER: PCT/IB02/00166
PRIOR FILING DATE: 2002-01-21
PRIOR APPLICATION NUMBER: US 10/050,902
PRIOR FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: US 60/396,635
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patentin version 3.2
SEQ ID NO 108
TENGTH: 14
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; OTHER INFORMATION: RANKL peptide EF loop
US-10-289-456-108
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US-10-105-232-267
; Sequence 267, Application US/10105232
; Publication No. US20030180328A1
; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 60/303,396
PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 08/198,139
PRIOR FILING DATE: 1994-02-17
NUMBER OF SEQ ID NOS: 535
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Best Local Similarity
Matches 5; Conserv
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APPLICANT: Maurer, Patrick
APPLICANT: Spohn, Gunther
                                                                                                                                                                                                                                                        APPLICANT: BOGOCH, ELENORE S:
TITLE OF INVENTION: REPLIKIN PEPTIDES IN RAPID REPLICATION OF GLIOMA CELLS
TITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS
FILE REFERENCE: 09425-46904
CURRENT APPLICATION NUMBER: US/10/105,232
CURRENT FILING DATE: 2002-03-26
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: BOGOCH
APPLICANT: BOGOCH
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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80.0%;
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Pred. No.
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Pred. No.
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RESULT 29
US-10-189-437-254 .
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; NAME/KEY: MOD_RES
; LOCATION: (5)
; OTHER INFORMATION: val or leu
US-10-105-232-267
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PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 09/146,755
PRIOR FILING DATE: 1998-09-04
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR APPLICATION NUMBER: 09/817,144
PRIOR FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: 08/198,139
PRIOR FILING DATE: 1994-02-17
                                                                                                                                                                      Sequence 254, Application US/10189437
Publication No. US20030194414A1
GENERAL INFORMATION:
APPLICANT: BOGOCH, SAMUEL
APPLICANT: BOGOCH, ELENORE S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local :
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APPLICANT: BOGOCH, ELENORE S.
TITLE OF INVENTION: REPLIKIN PEPTIDES AND ANTIBODIES THEREFORE
FILE REFERENCE: 09425/46905
CURRENT APPLICATION NUMBER: US/10/189,437
CURRENT FILING DATE: 2002-07-08
PRIOR APPLICATION NUMBER: 10/105,232
PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: 09/984,057
PRIOR APPLICATION NUMBER: 09/984,057
PRIOR FILING DATE: 2001-10-26
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ITLE OF INVENTION: REPLIKIN PEPTIDES IN RAPID REPLICATION OF GLIOMA CELLS
ITLE OF INVENTION: AND IN INFLUENZA EPIDEMICS
ILE REFERENCE: 09425-46904
URRENT APPLICATION NUMBER: US/10/105,232
URRENT FILING DATE: 2002-03-26
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b. US20030180328A1
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RESULT 30
US-10-189-437-257
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US-10-189-437-257
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PRIOR FILING DATE: 2002-03-26
PRIOR APPLICATION NUMBER: 09/984,057
PRIOR FILING DATE: 2001-10-26
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR APPLICATION NUMBER: 60/303,396
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR APPLICATION NUMBER: 60/278,761
PRIOR FILING DATE: 2001-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Pat
SEQ ID NO 257
LENGTH: 14
TYPE: PRT
              Sequence 16, Application US/09910552
Publication No. US20020197260A1
GENERAL INFORMATION:
APPLICANT: Granoff, Dan M.
APPLICANT: Moe, Gregory R.
TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT DEFINE UNIQUE
TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE PREPARATION OF VACCINE
FILE REFERENCE: 1238.002
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Matches
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LENGTH: 14
TYPE: PRT
ORGANISM: Influenza virus
FEATURE:
                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/910,552
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SOFTWARE: Patentt
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PRIOR FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: 60/278,761
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PRIOR FILING DATE: 2001-03-27
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TITLE OF INVENTION: REPLIKIN PEPTIDES AND ANTIBODIES THEREFORE
TILE REFERENCE: 09425/46905
URRENT APPLICATION NUMBER: US/10/189,437
URRENT FILING DATE: 2002-07-08
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LOCATION: (5)
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80.0%;
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Pred. No. 1e+03;
0; Mismatches
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APPLICANT: Moe, Gregory R.
TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT DEFINE UNIQUE
TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE PREPARATION OF VACCINE
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 1238.002
CURRENT APPLICATION NUMBER: US/10/643,465
CURRENT FILING DATE: 2003-08-19
PRIOR APPLICATION NUMBER: US/08/925,002
PRIOR FILING DATE: 1997-08-27
NUMBER OF SEQ ID NOS: 68
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
LENGTH: 8
TUDE OF SEQ ID NOS: 68
SOFTWARE: PATENTIN VER. 2.0
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PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 68
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
                                                                                                                                                                             Sequence 555, Application US/10117937
Publication No. US20030220239A1
GENERAL INFORMATION:
APPLICANT: CTL IMMUNO THERAPIES CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 16, Application US/10643465
Publication No. US20040077840A1
GENERAL INFORMATION:
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                APPLICANT: SIMARD, John, J.L.
APPLICANT: DIAMOND, David, C.
APPLICANT: LIU, Liping
APPLICANT: XIE, Zhidong
TITLE OF INVENTION: EPITOPE SEQUENCES
FILE REFERENCE: CTLIMM.027A
CURRENT APPLICATION NUMBER: US/10/117,937
CURRENT FILING DATE: 2002-04-04
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PRIOR APPLICATION NUMBER: US 60/282,211
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: sequence OTHER INFORMATION: a phage display peptide library
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ORGANISM: Artificial Sequence
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les 4; Conserv
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Pred. No. 1.2e+06;
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Pred. No. 1.
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; Sequence 126, Application US/09753126
; Patent No. US20020127219A1
; GENERAL INFORMATION:
; APPLICANT: OKKELS, JENS SIGURD
; APPLICANT: HALKIER, TORBEN
; APPLICANT: HALKIER, TORBEN
; APPLICANT: JENSEN, RIKKE BOLDING
; TITLE OF INVENTION: IMPROVED LYSOSOMAL ENZYMES ANI
; TITLE OF INVENTION: ACTIVATORS
; FILE REFERENCE: 31-000600US
; CURRENT APPLICATION NUMBER: US/09/753,126
; CURRENT FILING DATE: 2001-06-11
; PRIOR APPLICATION NUMBER: PA 1999 01891
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; LENGTH: 10
; TYPE: PRT
; ORGANISM: Homo Sapiens
US-09-780-053-472
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                                                                                                                                                                                                                                                          RESULT 35
US-09-753-126-126
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US-09-780-053-472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Rene S. Hubert
APPLICANT: Daniel B.H. Afar
APPLICANT: Pia M. Challita-Eid
APPLICANT: Mary Faris
APPLICANT: Tierin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/780,053
CURRENT FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/181,261
PRIOR FILING DATE: 2000-02-09
NUMBER OF SEQ ID NOS: 716
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 33.3
Conservative
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 555
                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Elana Levin
APPLICANT: Steve Chappell Mitchell
APPLICANT: Aya Jakobovits
TITLE OF INVENTION: 83P5G4: A TISSUE
TITLE OF INVENTION: HIGHLY EXPRESSED
FILE REFERENCE: 129.5USU1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: US 60/337,017
PRIOR FILING DATE: 2001-11-07
PRIOR APPLICATION NUMBER: US 60/363,210
PRIOR APPLICATION SERIOR 50/363,210
PRIOR FILING DATE: 2002-03-07
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                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
les 4; Conserv
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                                                                                                                                                                                                                                                                                                                                          GHQNSTF 10
                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        40.0%;
57.1%;
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Pred. No. 1.1e+03
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Pred. No. 1
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                                                                                                      AND LYSOSOMAL ENZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 10;
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FILING DATE:

1999-12-30

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COTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: peptide
US-09-753-126-126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 엉덩
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                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: peptide
US-09-896-896A-90
                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/217,497
PRIOR APPLICATION NUMBER: US 60/225,558
PRIOR APPLICATION NUMBER: US 60/225,558
PRIOR FILING DATE: 2000-08-16
PRIOR APPLICATION NUMBER: DK PA 2000 01027
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DK PA 2000 01092
PRIOR FILING DATE: 2000-07-14
PRIOR FILING DATE: 2000-07-14
PRIOR APPLICATION NUMBER: PCT/DK00/00743
PRIOR FILING DATE: 2000-12-29
PRIOR APPLICATION NUMBER: PCT/DK01/00090
PRIOR FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Pate
; SEQ ID NO 126
; TWOTH: 10
                                                                                      Best Loc
Matches
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Best Local Similarity 57.1
                                                                                                                                                                                                                                                                                                  SOFTWARE: P
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                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -09-896-896A-9C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: PEPTIDE EXTENDED GLYCOSYLATED POLYPEPTIDES FILE REFERENCE: 0217us210
CURRENT APPLICATION NUMBER: US/09/896,896A
CURRENT FILING DATE: 2001-06-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OR FILING DATE: 2000-06-12
OR APPLICATION NUMBER: 60/211,124
OR APPLICATION NUMBER: 60/211,124
IOR FILING DATE: 2000-06-12
IOR FILING DATE: 2000-06-30
IOR APPLICATION NUMBER: 60/217,497
IOR APPLICATION NUMBER: 60/217,497
IOR FILING DATE: 2000-07-11
IOR FILING DATE: 2000-07-11
                                                                                                          Local
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APPLICATION NUMBER: PA 200 00865
FILING DATE: 2000-06-02
APPLICATION NUMBER: 60/210,984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/174,652
                                                 2 YKNSKFH 8
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                                                                                      Similarity 57.: 4; Conservative
                                                                                                                                                                                                                                                                                                                          PatentIn Ver.
            YKSYKFY 10
                                                                                                       40.0%;
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                                                                                    Score 24; DB 10;
Pred. No. 1.1e+03;
2; Mismatches 1
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Pred. No. 1.1e+03
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                                                                                       Indels
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FILE REFERENCE: CTLIMM.027A
CURRENT APPLICATION NUMBER: US/10/117,937
CURRENT FILING DATE: 2002-04-04
PRIOR APPLICATION NUMBER: US 60/282,211
PRIOR FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: US 60/337,017
PRIOR FILING DATE: 2001-11-07
PRIOR APPLICATION NUMBER: US 60/363,210
PRIOR APPLICATION NUMBER: US 60/363,210
PRIOR FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 602
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 556
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                                                                                                                                                        ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-117-937-556
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US-10-117-937-556
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US-10-319-340-8
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 8, Application US/10319340
Publication No. US20030144211A1
GENERAL INFORMATION:
APPLICANT: Heavner, George A.
APPLICANT: McEver, Rodger P.
APPLICANT: Geng, Jian-Guo
TITLE OF INVENTION: Peptide Inhibitors of Inflammation Mediated by Selectins
FILE REFERENCE: CTC 102 CON DIV
CURRENT APPLICATION NUMBER: US/10/319,340
CURRENT FILING DATE: 2002-12-13
PRIOR APPLICATION NUMBER: 08/135,319
PRIOR APPLICATION NUMBER: 08/135,319
PRIOR FILING DATE: 1993-10-12
NUMBER OF CTO ID NOC: 26
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Best Local Similarity
Matches 4; Conserv
                                                                             Matches
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                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 556, A Publication No.
                                                                                                Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPLICANT: LIU, Liping
PPLICANT: XIE, Zhidong
ITLE OF INVENTION: EPITOPE SEQUENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Synthetic inhibitory peptide
                                                                                                                                                                                                                ENGTH: 10
                                     2 YKNSKFHRV 10
                                                                             Similarity
3; Conserv
YDNKQFEKI 10
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                                                                        40.0%; nilarity 33.3%; Conservative 3
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5. US20030220239A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 24; DB 14;
Pred. No. 1.1e+03;
1; Mismatches 0
                                                                                               Score 24; DB 15; Length 10; Pred. No. 1.1e+03;
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RESULT

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INFORMATION:

Application US/10330697 b. US20040009165A1

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CURRENT APPLICATION NUMBER: US/09/791,393
CURRENT FILING DATE: 2002-01-02
EARLIER APPLICATION NUMBER: GB 0004412.3
EARLIER FILING DATE: 2000-02-24
EARLIER APPLICATION NUMBER: GB 0030050.9
EARLIER FILING DATE: 2000-12-08
EARLIER APPLICATION NUMBER: US 60/254,830
EARLIER FILING DATE: 2000-12-12
NUMBER OF SEQ ID NOS: 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 1.
SOFTWARE: PatentIn Ver.
SEQ ID NO 126
                                                                                                                                                                                                                                                                                                                              Sequence 21, Application US/09791393
Publication No. US20030032200A1
GENERAL INFORMATION:
APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Rohlff, Christian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 31-000600US
CURRENT APPLICATION NUMBER: US/10/330,697
CURRENT FILING DATE: 2002-12-27
                                                                                                                                                                                                                                  TITLE OF INVENTION: Proteins, Genes and Their Use TITLE OF INVENTION: Diagnosis and Treatment of Big TITLE OF INVENTION: and Unipolar Depression FILE REFERENCE: 2543-1-001 N1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ITLE OF INVENTION: IMPROVED LYSOSOMAL ENZYMES AND LYSOSOMAL ENZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
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APPLICATION NUMBER: 60/174,652
FILING DATE: 2000-01-06
APPLICATION NUMBER: PA 200 00865
FILING DATE: 2000-06-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/210,984
FILING DATE: 2000-06-12
APPLICATION NUMBER: 60/211,124
FILING DATE: 2000-06-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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JENSEN, RIKKE BOLDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DATE: 2000-07-11
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57.1%;
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Pred. No.
                                                                                                                                                                                                                                                                                        Bipolar Affective Disorder (BAD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 10;
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APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri APPLICANT: Parekh, Rajesh Bhikhu

APPLICANT: Parekh, Christian

APPLICANT: Rohlff, Christian

APPLICANT: Terrett, Jonathan Alexander

APPLICANT: Tyson, Kerry Louise

TITLE OF INVENTION: Proteins, Genes and Their Use for INTILE OF INVENTION: Diagnosis and Treatment of Bipolar Affect ITILE OF INVENTION: and Unipolar Depression

FILE REFERENCE: 2543-1-001 N2

CURRENT APPLICATION NUMBER: US/09/791,389

CURRENT FILING DATE: 2001-02-23

PRIOR APPLICATION NUMBER: GB 0004412.3

PRIOR APPLICATION NUMBER: GB 0030050.9

PRIOR APPLICATION NUMBER: GB 030050.9

PRIOR APPLICATION NUMBER: US 60/254,830

PRIOR FILING DATE: 2000-12-08

PRIOR FILING DATE: 2000-12-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: PRT; ORGANISM: homo sapien US-09-791-389-21
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SOFTWARE: FastS
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Best Local Similarity 44.4%;
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                            Sequence 22, Application US/10264309
Publication No. US20040022794A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
INVENTION: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GYKNSKFHR 9
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                                                                                                                                                      HERATH, HERATH
KIMMEL, LIDA H.
PAREKH, RAJESH B.
POTTER, DAVID M.
ROHLFF, CHRISTIAN
SILBER, B. MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GYLNTGYQR 11
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                                                                                                                                                                                                                                                                                    FRIEDMAN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US20030032773A1
                                                                                ERLAND,
                                                      THOMAS R.
AND, P. TREY
D, ROBERT R.
                                                                                                                                                                                                                                                                                    . KATHRYN
, DAVID L.
                    STEPHEN A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 24;
Pred. No.
MOLECULES, POLYPEPTIDES AND USES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of Bipolar Affective Disorder (BAD)
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PRIOR APPLICATION NUMBER: 60/326,708; PRIOR FILING DATE: 2001-10-03; PRIOR FILING DATE: 2001-10-03; NUMBER OF SEQ ID NOS: 491; SOFTWARE: Patentin Version 2.1; SEQ ID NO 22; LENGTH: 11
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                                                                                                                                                                                                                       RESULT 44
US-09-734-002-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-264-309-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: synthetic peptide US-10-302-100B-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-302-100B-12
                                               Sequence 5, Application US/09734002
Patent No. US20010016333A1
GENERAL INFORMATION:
APPLICANT: Motoharu SEIKI et al.
TITLE OF INVENTION: NOVEL PROTEIN AND MONOCLONAL ANTIBODY SPECIFIC THERETO NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderoth, Lind & Ponack, L.L.P.
STREET: 2033 K Street, N.W., Suite 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 12 LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12, Application US/10302100B Publication No. US20040054131A1 GENERAL INFORMATION:
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Best Local (
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Best Local
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PRIOR FILING DATE: 1998-09-30
NUMBER OF SEQ ID NOS: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ballinger, Marcus
APPLICANT: Kavanaugh, Michael
TITLE OF INVENTION: Synthetic Peptides Having FGF Receptor
TITLE OF INVENTION: Affinity
FILE REFERENCE: 035784/263218
CURRENT APPLICATION NUMBER: US/10/302,100B
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: 09/407,687
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/264,309
CURRENT FILING DATE: 2002-10-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                       4;
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STATE: D.C.
COUNTRY: U.S.A.
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                                 Washington
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Pred. No. 1.
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Pred. No. 1.2e+03
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TOPOLOGY: Linear
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MOLECULE TYPE: Peptide
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SEQUENCE DESCRIPTION: SEQ
US-09-734-002-5
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                                                                                                                                 ; OTHER INFORMATION: BLyS binding polypeptide US-09-932-613-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-932-613-39
                                                                                                                                                               SOFTWARE: PatentIn version 3.1
SEQ ID NO 39
LENGTH: 14
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
APPLICANT: Beltzer, James P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 39, Application US/09932613
Publication No. US20030091565A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                               Best Local Similarity
                                                                                               Query Match
                                                                                                                                                                                                                                                                TITLE OF INVENTION: BINDING POLYPEPTIDES AND METHODS BASED THEREON FILE REFERENCE: Dyx-025.1 PCT; DYX-025.1 US CURRENT APPLICATION NUMBER: US/09/932,613 CURRENT FILING DATE: 2001-08-17 NUMBER OF SEQ ID NOS: 458
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/734,002

FILING DATE: 12-Dec-2000

CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/JP96/C1956

FILING DATE: July 12, 1996

ATTORNEY/AGENT INFORMATION:

NAME: Lee Cheng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-721-8200
TELEFAX: 202-721-8250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.5 inch, 1.44 mb
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
9
                              3 KNSKFH 8
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KHCKFH 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: Amino acid
STRANDEDNESS: Single
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REFERENCE/DOCKET NUMBER: <U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Wordperfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                    Beltzer, James P. Potter, M. Daniel
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                                                                  Conservative
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                                                                                40.0%;
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Pred. No. 1.6e+03;
1; Mismatches 0;
                                                                               Score 24; DB 10;
Pred. No. 1.6e+03;
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                                                                  Mismatches
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Search completed: August 30,

2004, 11:04:56

Job time : 14.4527 secs

Page 13

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Minimum DB seq length: 8
Maximum DB seq length: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
 Issued Patents AA:*
1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/iaa/pCTUS_COMB.pep:*
6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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Gapop 10.0 , Gapext 0.5
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60
 Copyright
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                                                                                                                                                                                                                                                                                                                      BB
US-08-481-88A-19
US-08-485-273A-19
US-08-973-559-19
US-08-973-553A-18
US-08-973-563A-24
US-08-973-563A-25
US-08-973-553A-27
US-08-973-559-18
US-08-973-559-27
US-08-973-559-27
US-08-973-559-27
US-08-973-559-27
US-08-973-559-27
US-08-973-559-36
US-08-973-559-36
US-08-973-559-36
US-08-973-559-36
US-08-973-559-36
US-08-993-235-4
US-08-993-235-4
US-08-993-235-6
US-08-993-235-7
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          sequence
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Sequence
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                                19, Appl
19, Appl
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18, Appl
18, Appl
18, Appl
24, Appl
25, Appl
27, Appl
36, Appl
36, Appl
36, Appl
37, Appl
38, Appl
39, Appl
39, Appl
39, Appl
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US-08-481-888A-19
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                                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Mili
                       STRANDEDNESS:
                                 : 11 amino amino amino acid
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             linear
  peptide
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| 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 25 | 25 | g | σ | 26.5 | O. |
| 0 | 40.0 | | | | 40.0 | 40.0 | 40.0 | 40.0 | 40.0 | 40.0 | 40.0 | | H | 4. | Α. | 44.2 | 4. |
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| 4 | N | N | 41 | 4. | μ | 4 | <u>ب</u> | 4 | ب | 4 | w | ហ | μ | N | N | \vdash | Ъ |
| US-08-475-399A-229 | -229 | 0-190-22 | | 87B- | 69-441 | 35-319A- | | US-09-417-608A-86 | US-08-215-805A-41 | 10-552-1 | 25-002-1 | PCT-US96-12632-8 | S-08-511-662- | 73-559-1 | US-08-973-563A-16 | US-08-441-914-6 | US-08-485-273A-16 |
| N.3 | equence | N) | · N) | ന | • | equence 8, | Sequence 5, Appli | equence 86, | \vdash | equence 16 | σ | equence 8 | equence 8, | equence | equence 16, Ap | App | Sequence 16, Appl |

ALIGNMENTS

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S-08-481-888A-19
Sequence 19, Application US/08481888A
Patent No. 5631228

GENERAL INFORMATION:
APPLICANT: Oppenheim, Frank G.
APPLICANT: Copenheim, Frank G.
APPLICANT: Copenheim, Frank G.
APPLICANT: No. Tao
APPLICANT: Roberts, F. Donald
TITLE OF INVENTION: HISTATIN-BASED PEPTIDES
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
COUNTRY: US
ZIF: O2173
COMPUTER READABLE FORM:
COUNTRY: HEXINGTON
STATE: MA
COMPUTER FRADABLE FORM:
COMPUTER SEADABLE FORM:
COUNTRY: US
ZIF: O2173
COMPUTER SEADABLE
COUN
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US-08-485-273A-19
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                              Matches
                                                               Query Match
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GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/287,717
FILING DATE: 09-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/145,030
FILING DATE: 28-OCT-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/145,030
                                                                                                                                                                                                                                                                      TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 07/786,571
FILING DATE: 01-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 617-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                               Local Similarity
les 6; Conserv
                                                                                                            OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
                                                                                                                                           NAME/KEY: Region LOCATION: 1..11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/485,273A FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TLE OF INVENTION: D-Amino Acid His TLE OF INVENTION: Anti-Fungal and
                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
  1 GYKNSKFH 8
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                                                                                                                                                                                                                                       amino acid
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                                                                                                                                                                                                                                                         11 amino acids
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Spacciapoli, Peter
                                                                                                                                                                                                           linear
                                                                                                                                                                                         peptide
                                                                                                                                                                                                                                                                                                                                                                                   David E.
                                                44.2%;
75.0%;
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75.0%;
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                                                                                                            /note= "At least one amino acid
must have a D configuration."
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                                                Score 26.5;
Pred. No. 1e
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Pred. No. 16
                               Mismatches
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1e+02;
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                                                               Length 11;
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                               Indels
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NAME/KEY: Region
; LOCATION: 1..11
; OTHER INFORMATION:
; OTHER INFORMATION:
US-08-973-563A-19
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                                                                                      RESULT 4
US-08-973-559-19
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US-08-973-563A-19
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Sequence 19, Application US/08973559
Patent No. 5912230
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: XU, TAO
                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                      Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 07-JUN-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 78
INFORMATION FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,56
                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Oppenhei
APPLICANT: Xu, Tao
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PEI
                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PPLICANT: Friden, Philip M.
TTLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based
TLE OF INVENTION: Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
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6; Conserva
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                                                                                                                                                        5 GYKR-KFH 11
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                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                      peptide
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75.0%;
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                                                                                                                                                                                                                                                     Patent No. 5631228
GENERAL INFORMATION:
APPLICANT: Oppenheim, Frank G.
APPLICANT: Xu, Tao
APPLICANT: Roberts, F. Donald
TITLE OF INVENTION: ANTI-FUNGAL
TITLE OF INVENTION: HISTATIN-BAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 781-861-6240
TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: peptide -08-973-559-19
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APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-01A2
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 781-861-6240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                    TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/973,559
FILING DATE: 07-JUN-1996
CLASSIFICATION: 514
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                                                                                                  COUNTRY: U
ZIP: 02173
                                                                                                                                                      CITY: Lexington
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Similarity 75.0%;
6; Conservative
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                                                                                                                                                                              Two Militia Drive
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                                                                                                                                                                                            Hamilton, Brook, Smith & Reynolds, P.C
                                                                                                                                                                                                                                  ANTI-FUNGAL AND ANTI-BACTERIAL HISTATIN-BASED PEPTIDES: 23
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RESULT 6
US-08-485-273A-18
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Best Local Similarity 75.0
Thehes 6; Conservative
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APPLICANT: Oppenhe
APPLICANT: Xu, Tao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,273A
FILING DATE: 07-JUN-1995
                                                                 CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/287,717
FILING DATE: 09-AUG-1994
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PE
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/786,571
FILING DATE: 01-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Spacciapoli, Peter
TITLE OF INVENTION: D-Amino Acid Histatin-Based Peptides as
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial Agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                      STREET: Two Mil:
CITY: Lexington
STATE: MA
APPLICATION NUMBER: US 08/145,030 FILING DATE: 28-OCT-1993 RIOR APPLICATION DATA: US 07/786,571 APPLICATION NUMBER: US 07/786,571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/287,717 FILING DATE: 09-AUG-1994
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                                                                                                                                                                                                                                                                                                                                COUNTRY: UZIP: 02173
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US 08/145,030
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75.0%;
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Pred. No. 1.1e
0; Mismatches
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.1e+02;
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Best Local Similarity 75.0
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                                                                                     CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-02A2
TELEFAX: 781-861-9540 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                          ZIP: 02173

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,563A
FILING DATE: 07-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-02A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
                                                    TELEPHONE: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Oppenhei
APPLICANT: Xu, Tao
APPLICANT: Spaccia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Spacciapoli, Peter
APPLICANT: Roberts, F. D.
APPLICANT: Friden, Philip M.
TITLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based
TITLE OF INVENTION: Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Region
LOCATION: 1..12
OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1wc
STREET: Lexington
                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: UZIP: 02173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
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Two Militia Drive
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75.0%;
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must have a D configuration."
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OTHER INFORMATION:
OTHER INFORMATION:
US-08-973-563A-18
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Best Local Similarity
"^+ches 6; Conserva
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                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/485,273

FILING DATE: 07-JUN-1995

ATTORNEY/AGENT INFORMATION:

NAME: Brook, David E.

REGISTRATION NUMBER: 22,592

REFERENCE/DOCKET NUMBER: PER95-02A2

TELECOMMUNICATION INFORMATION:

TELEPHONE: 781-861-6240
                                                                                                                                            TELEFAX: 781-801
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: TRNGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: PATEN: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Ver.

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/973,563A

FILING DATE: 07-JUN-1996

CLASSIFICATION: 514
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APPLICANT: OppenhappLICANT: Xu, Tax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
NAME/KEY: Region LOCATION: 1..12 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: Two Milit
CITY: Lexington
STATE: MA
COUNTRY: US
ZIP: 02173
                                                                                       MOLECULE TYPE: peptide
                                                                                                         TYPE: amino acid STRANDEDNESS: TOPOLOGY: linear
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Friden, Philip M.
FVENTION: Anti-Fungal D-Amino Acid Histatin-Based
NVENTION: Peptides
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75.0%;
  /note= "At least one amino acid
must have a D configuration."
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Pred. No. 1.1e+02;
0; Mismatches 1;
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Query Match
Best Local Similarity
Matches 6; Conserv

Conservative

44.2%; 75.0%;

Score 26.5; Pred. No. 1

1.1e+02; DB 2;

Length 12;

Indels

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Gaps

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; NAME/KEY: Region
; LOCATION: 1..12
; OTHER INFORMATION:
; OTHER INFORMATION:
US-08-973-563A-25
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                                                                                                                                            Matches
                                                                                                                                                                                Query Match
Best Local :
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SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 781-861-6240
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,9
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 3'CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Lexington
STATE: MA
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FILING DATE: 07-JUN-1996
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                                                       1 GYKNSKFH 8
                                                                                                                                       Similarity 6; Conserv
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GYKR-KFH 12
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Roberts, F. D.
Friden, Philip M.
Friden, F. D.
Friden, Philip M.
Frid
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Xu, Tao
                                                                                                                                            Conservative
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                                                                                                                                                                       44.2%;
                                                                                                                                                                                                                                                                                                              /note= "At least one amino acid
must have a D configuration."
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                                                                                                                                          0; Mismatches
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                                                                                                                                                                    Score 26.5; DB 2;
Pred. No. 1.1e+02;
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            Sequence 36, Application US/08973563A Patent No. 5885965
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Patent No. 5
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TELEFAX: 781-861-9540
:NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 781-861-6240
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                                                                                                                                                                                                                                                             LOCATION: 1..12
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                              Local Similarity les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Brook, David E.
REGISTRATION NUMBER: 22,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/973,563A FILING DATE: 07-JUN-1996 CLASSIFICATION: 514
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INFORMATION:
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                                                                                                                                                1 GYKNSKFH 8
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Friden, Philip M.
VENTION: Anti-Fungal D-Amino Acid Histatin-Based
                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                    linear
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must have a D configuration."
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                                                                                                                                                                                                Score 26.5;
Pred. No. 1
                                                                                                                                                                                                1.1e+02;
                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                               Length 12;
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1: Roberts, F. D.
1: Friden, Philip M.
1: Friden, Philip M.
1: INVENTION: Anti-Fungal D-Amino Acid Histatin-Based
1: INVENTION: Peptides
7: 75775875; 37

Oppenheim, Frank G.

<u>,,</u>

Spacciapoli, Peter

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INFORMATION FOR SEQ ID NO: 36
SEQUENCE CHARACTERISTICS:
FRACTH: 12 amino acids
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Best Local (
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,273
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: BYOCK DAVIG E
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                       ent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                   TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  44.2%;
Local Similarity 75.0%;
les 6; Conservative
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                                                                    COUNTRY: US
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                                                                                                                                   STREET:
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                                                                                                            SSEE: Hamilton, Brook, Smith & Reynolds, P.C.
F: Two Militia Drive
Lexington
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                                                                                                   ΜA
                                                                                                                                                                                                                                                                                                      OPPENHEIM, FRANK G
                                                                                                                                                                                                                                    SPACCIAPOLI, PETE FRIDEN, PHILIP M.
                                                                                                                                                                                                                                                                       ROBERTS, F. D.
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                                                                                                                                                                                      Anti-Fungal and Anti-Bacterial Histatin-Based Peptides : 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "At least one amino acid
must have a D configuration."
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RESULT 13
US-08-973-559-24
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Best Local Similarity 75.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-01A2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 781-861-6240
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: peptide
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APPLICATION NUMBER: US 0
FILING DATE: 07-JUN-1995
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: PE
                                                                                                                                                                                                                      APPLICATION NUMBER:
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SPACCIAPOLI, PETER
FRIDEN, PHILIP M.
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07-JUN-1996
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                                                                                                                                                                                                                                                    Release #1.0, Version #1.30
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Query Match
Best Local Similarity
Watches 6; Conserve
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US-08-973-559-25
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Best Local Similarity 75.0%;
Matches 6; Conservative
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,559
FILING DATE: 07-JUN-1996
CLASSIFICATION: 514
                                                                                                                                                                                                                        TELEFAX: 781-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT: OPPENH
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LENGTH: 12 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: SPACCIAPOLI, PETER
APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                              MOLECULE TYPE: peptide
-973-559-25
                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 781-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C STREET: Two Militia Drive
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                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                            TYPE: amino acid
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 GYKR-KFH 12
                                GYKNSKFH
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                                                           44.2%;
nilarity 75.0%;
Conservative
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XU, TAO
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Pred. No. 1.
                                                               Score 26.5; DB 2; Length 12; Pred. No. 1.1e+02; o; Mismatches 1; Indels
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GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FF
APPLICANT: XU, TAO
APPLICANT: ROBERTS, F. I
APPLICANT: SPACCIAPOLI,

OPPENHEIM, FRANK G.

F.

PETER

APPLICANT: FRIDEN, PHILIP M.
APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
NUMBER OF SEQUENCES: 37

ADDRESS: Hamilton, Brook, Smith & Reynolds, P.C.

Patent No.

Application US/08973559

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Query Match
Best Local Similarity
Marches 6; Conserva
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US-08-973-559-27
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RESULT 16
US-08-973-559-36
                                                                                                                                                                                                                                                                                                                        ATTOKNES,....

NAME: Brook, David E.

NAME: Brook, David E.

REGISTRATION NUMBER: 22,592

REFERENCE/DOCKET NUMBER: PER9

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION 381-861-6240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: XU, TAO
                                                                                                                                                                                                                                                                                                          TELEFAX: 781-8
                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0,

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: ROBERTS, F. D.
APPLICANT: SPACCIAPOLI, PETER
APPLICANT: FRIDEN, PHILIP M.
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial
TITLE OF INVENTION: Histatin-Based Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
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CITY: Lexington
STATE: MA
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TOPOLOGY: li
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                                                             GYKR-KFH 12
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                                                                                                                             Conservative
                                                                                                                                                                                                                            linear
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07-JUN-1996
                                                                                                                                            44.2%;
75.0%;
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                                                                                                                                           Score 26.5;
Pred. No. 1.
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                                                                                                                                                           Length 12;
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RESULT 18
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                                                                                                                                                                                   LENGTH: 12
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-993-235-4
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                                                                                                                                                                                                                                                   APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. DONALD
APPLICANT: ROBERTS, F. DONALD
APPLICANT: XU, TAO
APPLICANT: SPACCIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT APPLICATION NUMBER: US/08/993,235
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4
                                                                                                                     Matches
                                                                                                                                   Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/08993235 Patent No. 6084064
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Best Local &
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APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-01A2
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 101-06140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 36:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/973,559 FILING DATE: 07-JUN-1996
                                                                                                                                    Local Similarity
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CITY: Lexington
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es 6; Conserv
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                                                                                   1 GYKNSKFH 8
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                                                                                                                     Conservative
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                                                                                                                                   44.2%;
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                                                                                                                   Score 26.5; DB 3;
Pred. No. 1.1e+02;
0; Mismatches 1;
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Pred. No. 1.1e+02;
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APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. DONALD
APPLICANT: XU, TAO
APPLICANT: SPACCIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT APPLICATION NUMBER: US/08/993,235
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
LENGTH: 12
TYPE: PRT
CRGANISM: Homo sapiens
US-08-993-235-7
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US-09-226-666-1
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; ORGANISM: Homo sapiens
US-08-993-235-5
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                 GENERAL INFORMATION:
APPLICANT: Spacciapoli, Peter
APPLICANT: Spacciapoli, David M.
APPLICANT: Friden, David M.
APPLICANT: Friden, PATHIND M.
TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
CURRENT APPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 18
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Best Local Similarity
Matches 6; Conserv
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Patent No. 6084064
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 5
LENGTH: 12
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Patent No. 6084064
GENERAL INFORMATION:
                                                                                                                                                                                     Sequence 1, Application US/09226666A Patent No. 6528488
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APPLICANT: ROBERTS, F. DONALD
APPLICANT: XU, TAO
APPLICANT: SPACCIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
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CURRENT APPLICATION NUMBER: US/08/993,235
CURRENT FILING DATE: 1997-12-18
SOFTWARE: FastSEQ for Windows Version 4.0
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Local Similarity 75.0%;
les 6; Conservative
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75.0%;
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Pred. No. 1.1e+02;
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; Sequence 16, Application US/09226666A
; Patent No. 6528488
; GENERAL INFORMATION:
; APPLICANT: Spacciapoli, Peter
APPLICANT: Friden, David M.
; APPLICANT: Friden, Phillip M.
; APPLICANT: Friden, Phillip FOR TREATING CYSTIC FIBROSIS
; TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
; FILE REFERENCE: 50032/007001
; CURRENT APPLICATION NUMBER: US/09/226,666A
; CURRENT FILING DATE: 1999-01-08
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 16
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                                                                  ; OTHER INFORMATION: Derived from Homo sapiens US-09-226-666-16
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Best Local S
Matches 6
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APPLICANT: Rothstein, David M.
APPLICANT: Rothstein, David M.
APPLICANT: Friden, Phillip M.
TITLE OF INVENTION: METHOD FOR TREATING CYSTIC FIBROSIS
FILE REFERENCE: 50032/007001
CURRENT APPLICATION NUMBER: US/09/226,666A
CURRENT FILING DATE: 1999-01-08
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13
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Patent No. 6528488
GENERAL INFORMATION:
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Best Local Similarity
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 Query Match
Best Local Similarity 75.0%;
Matches 6; Conservative
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ORGANISM: Artificial Sequence
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LENGTH: 12
                                                                                                            TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                     FEATURE:
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Score 26.5; DB 4;
Pred. No. 1.1e+02;
0; Mismatches 1;
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Pred. No. 1.1e+02;
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GENERAL INFORMATION:

APPLICANT: OPPENHEIM, FRANK G.

APPLICANT: ROBERTS, F. DONALD

APPLICANT: XU, TAO

APPLICANT: SPACCIAPOLI, PETER

TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES

FILE REFERENCE: 50032/002001

CURRENT APPLICATION NUMBER: US/08/993,235

CURRENT FILING DATE: 1997-12-18

NUMBER OF SEQ ID NOS: 13

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 4
                                                                                                                                                                                                                     APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: XU, TAO
APPLICANT: XU, TAO
APPLICANT: XU, TAO
APPLICANT: SPACCIAPOLI, PETER
TITLE OF INVENTION: ANTIFUNGAL AND ANTIBACTERIAL PEPTIDES
FILE REFERENCE: 50032/002001
CURRENT APPLICATION NUMBER: US/08/993,235
CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 5
LENGTH: 12
TYPE: PRT
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-993-235-4
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US-08-993-235-4
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RESULT 25
US-08-993-235-7
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                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                              Matches
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Pred. No. 1.1e+(
0; Mismatches
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Sequence 7, Application US/08993235
Patent No. 6531573
GENERAL INFORMATION:
APPLICANT: OPPENHEIM, FRANK G.
APPLICANT: ROBERTS, F. DONALD

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US-08-287-717-6
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-993-235-7
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CURRENT FILING DATE: 1997-12-18
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
 Query Match 44.2%;
Best Local Similarity 75.0%;
Matches 6; Conservative
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Best Local Similarity
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Patent No. 5486503
                                                                                                                  TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                              TOPOLOGY: li
                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 0 FILING DATE: 01-NOV-1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                 TELEPHONE: 617-861-6240
                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: I
FILING DATE:
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                                                                                                                                                                                                                                NAME: Granahan, Patricia REGISTRATION NUMBER: 32, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LICANT: Xu, Tao
LE OF INVENTION:
LE OF INVENTION:
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)GY: linear
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                                                                                peptide
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                                                                                                                                                                                                                                                                                                                  US 07/786,571
                                                                                                                                                                                                                                                                                                                                                 US/08/145,030
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                                                                                                                                                                                                                                              32,227
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Score 26.5; DB 1;
Pred. No. 1.3e+02;
0; Mismatches 1;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4;
                                Length 14;
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 Gaps
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                                                         US-08-485-273A-16
                                                                          RESULT 28
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US-08-481-888A-16
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Sequence 16, Application Patent No. 5646119
GENERAL INFORMATION:
                                                                                                                                                                                                          Best Loc
Matches
                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/481,888A FILING DATE: 07-JUN-1995 CLASSIFICATION: 514 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/287,717 FILING DATE: 09-AUG-1994 PRIOR APPLICATION NUMBER: US 08/145,030 APPLICATION NUMBER: US 08/145,030 FILING DATE: 28-OCT-1993 PRIOR APPLICATION NUMBER: US 07/786,571 APPLICATION NUMBER: US 07/786,571 APPLICATION NUMBER: US 07/786,571 FILING DATE: 01-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 14 amino acidi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
ORBEITON NUMBER. 115/06/461 8888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reymolds, P.C.
STREET: Two Militia Drive
CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                                                                  TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Roberts, F. Donald
TITLE OF INVENTION: ANTI-FUNGAL ANI
TITLE OF INVENTION: HISTATIN-BASED
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: UZIP: 02173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
                                                                                                                                                                     1 GYKNSKFH 8
                                                                                                                                                                                                          Similarity 6; Conserv
                                                                                                                                    GYKR-KFH
                                                                                                                                                                                                                                                                                                                                                       amino acid
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                                                                                                                                                                                                                                                                                                                                                                           14 amino acids
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                                                                                                                                                                                                                                                                                                                      linear
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                                                                                                                                                                                                                         44.2%;
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                                         US/08485273A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22,592
                                                                                                                                                                                                        Score 26.5; DB 1;
Pred. No. 1.3e+02;
0; Mismatches 1;
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                                                                                                                                                                                                                                          Length 14;
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Oppenheim, Xu, Tao

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US-08-441-914-6
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                           Sequence 6, Applicati
Patent No. 5696078
GENERAL INFORMATION:
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FILING DATE: 01-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: PER95-02A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
                                                         APPLICANT: Oppenheim, FI
APPLICANT: Xu, Tao
TITLE OF INVENTION: No.
TITLE OF INVENTION: Thei
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE: Region LOCATION: 1..14 OTHER INFORMATION: 0..14 OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 617-861-9540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
LENGTH: 14 amino acids
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/145,030
FILING DATE: 28-OCT-1993
PRIOR APPLICATION DATA:
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ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Spacciapoli, Peter
TITLE OF INVENTION: D-Amino Acid Histatin-Based Peptides as
TITLE OF INVENTION: Anti-Fungal and Anti-Bacterial Agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
ZIP: 02173
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TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
                                          DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                             1 GYKNSKFH 8
SEE: Hamilton, Brook, Smith & Reynolds, P.C.
: Two Militia Drive
Lexington
                                                                                                                                                                                                                  Application US/08441914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14 amino acids
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                                                                                                                                                    Oppenheim, Frank G.
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                                                                                                   No. 5696078el Anti-Fungal Peptides and Uses Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                    44.28;
75.08;
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must have a D configuration."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 30
US-08-973-563A-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: peptide US-08-441-914-6
                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Oppenhe:
APPLICANT: Xu, Tao
APPLICANT: Spaccial
APPLICANT: Roberts
                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/145,030
FILING DATE: 28-OCT-93
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/786,571
APPLICATION NUMBER: 07/786,571
ATTORNEY/AGENT INFORMATION:
NAME: David E. Brook
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Veri
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/973,563A
FILING DATE: 07-JUN-1996
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/287,717
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                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                        ADDRESSEE: Hamili
STREET: Two Milit
CITY: Lexington
STATE: MA
                                                                                                                                                                                                                                                              NPPLICANT: Roberts, F. D.
NPPLICANT: Friden, Philip M.
IITLE OF INVENTION: Anti-Fungal D-Amino Acid Histatin-Based
IITLE OF INVENTION: Peptides
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: M
                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 22,592
REFERENCE/DOCKET NUMBER: BU91-17F2Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GYKR-KFH 7
                                                                                                                                         02173
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                                                                                                                                                                                                            E: Hamilton, Brook,
Two Militia Drive
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75.0%;
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Pred. No. 1
                                                                                                                                                                                                                             Smith & Reynolds,
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                                                      Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                       #1.30
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FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/481,888
FILING DATE: 07-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
REGISTRATION NUMBER: PER95-01A
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TOTREPHONE: 781-861-6240
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US-08-973-559-16
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Best Local S
Matches 6
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Patent No. 5912230
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INFORMATION FOR SEQ ID NO: 16:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reymolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: PEI
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FITLE OF INVENTION:
FITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: OPPENHEIM, FRANK G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/973,559
FILING DATE: 07-JUN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-Fungal and Anti-Bacterial
Histatin-Based Peptides
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US-08-511-662-8
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Best Local Similarity 75.0%;
Matches 6; Conservative
                                                             Matches
                                                                                           Query Match
                                                                              Best Local Similarity
                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 14 amino acids
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                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: UTS
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION: 515/418-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
Tom DC Com
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APPLICANT: Hughes, Jr., 11.
APPLICANT: Smith, Eric M.
Compo
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NAME: Hodgins, Daniel S.
REGISTRATION NUMBER: 31,
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ADDRESSEE: Arnold, Wh.
STREET: P.O. Box 4433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Compositions for Conferring Immunogenicity TITLE OF INVENTION: to a Substance and Uses Thereof NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                           TELEPHONE: 515/---
TELEPHONE: 515/---
TELEPHONE: 515/474-7577
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: Concur CLASSIFICATION: 530
                                                                                                                                                                     STRANDEDNESS:
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 5 YLNLRFH 11
                              2 YKNSKFH 8
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                                                              4; Conservative
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                                                                             41.7%;
57.1%;
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Pred. No. 1.3e+02;
0; Mismatches 1
                                                                             Score 25; DB 1;
Pred. No. 2e+02;
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                                                                                              DB 1;
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                                                                                             Length 12;
                                                               Indels
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RESULT 33 PCT-US96-12632-8

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RESULT 34
US-08-925-002-16
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INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 4
                                                                                                     Sequence 16, Application US/08925002 Patent No. 6048527 GENERAL INFORMATION:
APPLICANT: Granoff, Dan M.
APPLICANT: Moe, Gregory R.
TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT DEFINE UNIQUE
TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE PREPARATION OF VACCINE
TITLE OF INVENTION: COMPOSITIONS
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APPLICANT: TELEFAX: (512) 499-4020

APPLICANT: TELEFAX: (512) 499-4020

TITLE OF INVENTION: COMPOSITIONS FOR CONFERRING

TITLE OF INVENTION: IMMUNOGENICITY TO A SUBSTANCE AND

TITLE OF INVENTION: USES THEREOF

TITLE OF SECUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 77210-4433
ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS/ASCII
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
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REFERENCE/DOCKET NUMBER: UTILELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/511,662
FILING DATE: 04 August 1995
CLASSIFICATION:
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ADDRESSEE: Arnold, Wh
STREET: P.O. Box 4433
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NAME: HODGINS, DANIEL S.
REGISTRATION NUMBER: 31,
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TELEPHONE: 512/474-7577
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                                                                                                                                                                                                                                                                                                                                       Local Similarity
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POSTAL CODE:
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                                                                                                                                                                                                                                                                                                                    Score 25; DB
Pred. No. 2e+0
1; Mismatches
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2e+02;
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Sequence 41, Application US/08215805A Patent No. 5559008

GENERAL INFORMATION:
APPLICANT: Chang, Yung-Fu
TITLE OF INVENTION: LEUKOTOXIN GENE FROM PASTEURELLA
TITLE OF INVENTION: SUIS
NUMBER OF SEQUENCES: 84
CORRESPONDENCE ADDRESS:

STATE:
COUNTRY: UC.
14603

New York Y: USA

ADDRESSEE: Nixon, Hargr: STREET: Clinton Square, CITY: Rochester

Hargrave, Devans & Doyle quare, P.O. Box 1051

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TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 1238.002
CURRENT APPLICATION NUMBER: US/09/910,552
CURRENT FILING DATE: 2001-07-23
PRIOR APPLICATION NUMBER: 09/494,822
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 68
SOFTWARE: Patentin Ver. 2.0
SEQ ID NOS: 68
TYPE
                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: sequence from ; OTHER INFORMATION: a phage display peptide library US-09-910-552-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: sequence from ; OTHER INFORMATION: a phage display peptide library US-08-925-002-16
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RESULT 36
US-08-215-805A-41
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CURRENT FILING DATE: 1997-08-27
NUMBER OF SEQ ID NOS: 68
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
LENGTH: 8
                                                                                                                                      Matches
                                                                                                                                                    Query Match
Best Local
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Best Local (
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                                                                                                                                   Similarity 4; Conserv
                                                                   GYEVQPFH
                                                                                                  GYKNSKFH
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Pred. No. 3e+0
1; Mismatches
                                                                                                                                   Score 24; DB 4;
Pred. No. 3e+05;
1; Mismatches
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3e+05;
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RESULT 38
US-07-909-122-5
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Best Local S
Matches 4
           Sequence 5, Application US/07909122 Patent No. 5415995 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.2
SEQ ID NO 86
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 86,
                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Roggen, Erwin
APPLICANT: Ernst, Steffen
TITLE OF INVENTION: Low Allergenic Protein Variants
FILE REFERENCE: 5676.200-US
CURRENT APPLICATION NUMBER: US/09/417,608A
CURRENT FILING DATE: 1999-10-13
NUMBER OF SEQ ID NOS: 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -09-417-608A-86
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 22-MAR-199
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                            ORGANISM: Artificial FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            itent No.
                                                                                                                                                                                                                                                             OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPLICANT: Olsen, Arne
 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORIGINAL SOURCE: ORGANISM: Actinobacillus pleuropneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 34,103
REFERENCE/DOCKET NUMBER: 19603/61 (D-1329A)
TELECOMMUNICATION INFORMATION:
TELEPHONE: (716) 263-1636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                       Local Similarity 66.7%; es 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
                                                                                                                                                  YKNSKF 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/09417608A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 amino acids
SCHOOLNIK, GARY K.
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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Pred. No. 3e+05;
2; Mismatches
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; TOPOLOGY: linear US-07-909-122-5
                                                                                                                          NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
SEQ ID NO 8
LENGTH: 10
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
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                                                                                         , OTHER INFORMATION: Synthetic inhibitory peptide US-08-135-319A-8
                                                                                                                                                                                                                                                                                                                                                                                                       US-08-135-319A-8
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Best Local (
                                                            Query Match
                               Matches
                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                              equence 8, Application US/08135319A
                                                                                                                                                                                                                                      APPLICANT: Heavner, George A.
APPLICANT: McEver, Rodger P.
APPLICANT: Geng, Jian-Guo
TITLE OF INVENTION: Peptide Inhibitors of Inflammation Mediated by Selectins
FILE REFERENCE: CTC 102 CON
CURRENT APPLICATION NUMBER: US/08/135,319A
CURRENT FILING DATE: 1993-10-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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NAME: BENZ, WILLIAM H.
REGISTRATION NUMBER: 25,952
REFERENCE/DOCKET NUMBER: 2860
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
                                                                                                                                                                                                                                                                                                                                       PPLICANT: Heavner, George
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                              Local
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2 YKNSK 6
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                              Similarity 80.
4; Conservative
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3; Conserv
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milarity 37.5%;
Conservative
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VENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA
VENTION: VIRUS
                                            40.0%;
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                              Score 24; DB 4; L. Pred. No. 2.6e+02; 1; Mismatches 0;
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Pred. No. 2.6e
2; Mismatches
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                                                            DB 4; Length 10;
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                                  Indels
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Sequence 88, Application US/09446787B; Patent No. 6541198; GENERAL INFORMATION:
APPLICANT: Akzo No. 6541198el N.V.
APPLICANT: Yan Kesel-Koenis, Marjolijin J.
APPLICANT: Van Kesel-Koenis, Marjolijin J.
TITLE OF INVENTION: Antibodies and other bind
TITLE OF INVENTION: antigens
FILE REFERENCE: 9310-19
CURRENT APPLICATION NUMBER: US/09/446,787B; CURRENT FILING DATE: 2000-01-13
NUMBER OF SEQ ID NOS: 97
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US-09-446-787B-88
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US-08-269-441A-3
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Best Local S
Matches 3
SOFTWARE: Pa
SEQ ID NO 88
LENGTH: 12
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/269,441A
FILING DATE: 30-JUN-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Haile Ph.D. Lisa A.
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07257/009001
TELECOMMUNICATION INFORMATION:
TELEPAX: (619) 678-5070
TELEFAX: (619) 678-5070
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APPLICANT: Rearden, Ann
TITLE OF INVENTION: A NOVEL AUTOANTIGEN, PINCH
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
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LENGTH: 11 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
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LOCATION:
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TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                             1 FKNDPYH 7
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                                                                                                                                                                                                                                                                                                                                                                                                             h 40.0%;
Similarity 42.9%;
3; Conservative
                                   PatentIn version 3.1
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1..11
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linear
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Pred. No. 2.8e+02;
2; Mismatches 2;
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TYPE: PRT
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RESULT 43
US-08-480-190-229
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US-09-407-687-12
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EARLIER FILING DATE: 1998-09-30
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 12
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                    Patent No. 5827516
GENERAL INFORMATION:
APPLICANT: Robert G. Urban
APPLICANT: Roman M. Chicz
APPLICANT: Dario A. A. Vignali
APPLICANT: Mary L. Hedley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity 30.1
Matches 3; Conservative
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APPLICANT: Ballinger, Marcus
APPLICANT: Kavanaugh, Michael
TITLE OF INVENTION: Synthetic Peptides Having FGF Receptor
TITLE OF INVENTION: Affinity
FILE REFERENCE: 1517.001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/407,687 CURRENT FILING DATE: 1999-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: synthetic peptide
              ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                 APPLICANT: Lawrence J. Stern
APPLICANT: Jack L. Strominger
TITLE OF INVENTION: IMMUNOMODULATORY PEPTIDES
                                                                                                                                                                 STREET: 225 Franklin CITY: Boston STATE: Massachusetts
                                                                                                                               COUNTRY: U.S.A. ZIP: 02110-2804
                                                                                                                                                                                            ADDRESSEE: Fish & ALCOHOLISTREET: 225 Franklin Street
 APPLICATION NUMBER:
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US/08/480,190
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Pred. No. 3.1e+02;
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Pred. No. 3.1e+02;
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Best Local Similarity
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/077,255
FILING DATE: June 15, 1993
APPLICATION NUMBER: 07/925,460
FILING DATE: AUGUST 11, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00246/168001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-8906
TELEPAX: (617) 542-8906
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: 08/077,255
FILING DATE: June 15, 1993
APPLICATION NUMBER: 07/925,460
FILING DATE: August 11, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                            ZIP: 02110-2804

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

COMPUTER: IBM PS/2 Model 50Z or 55SX

OPERATING SYSTEM: MS-DOS (Version 5.0)

SOFTWARE: WordPerfect (Version 5.1)

CURRENT APPLICATION DATA:
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NAME: Clark, Paul 1.

NAME: 30,162

REGISTRATION NUMBER: 30,162

REFERENCE/DOCKET NUMBER: 00246/168001

TELECOMMUNICATION INFORMATION:

THE COMMUNICATION 542-5070

THE CHORNE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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E: Fish & Richardson
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Dario A. A. Vignali
Mary L. Hedley
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jack L. Strominger
VENTION: IMMUNOMODULATORY PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lawrence J. Stern
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Robert G. Urban
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       June 15, 1993
WUMBER: 07/925,460
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Pred. No. 3.4e+02;
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US-08-488-379-229
                                                                                                                            US-08-475-399A-229
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 40.0%;
Best Local Similarity 44.4%;
Matches 4; Conservative
                                                            Matches
                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/475,399A
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/077,255
FILING DATE: 15-JUN-1993
APPLICATION NUMBER: 07/925,460
FILING DATE: 11-AUG-1992
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00246/168003
TELEPHONE: 617/542-507
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 13
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 13 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Vignali, Dario A.A.
APPLICANT: Hedley, Mary L.
APPLICANT: Stern, Lawrence J.
APPLICANT: Strominger, Jack L.
TITLE OF INVENTION: IMMUNOMODULATORY PEPTIDES
NUMBER OF SEQUENCES: 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino STRANDEDNESS
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225 Franklin Street
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Pred. No. 3.4e+02;
2; Mismatches 3
                                                             Score 24; DB 4;
Pred. No. 3.4e+02;
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                                                                                          Length 13
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Perfect score:
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| | 8mh51 homo | 8mh30 homo | 8mh31 homo s | 8mh14 homo s | 8mh56 homo | 8mh38 homo s | 8mh16 homo : | 5 homo s | 7 homo | 0 | 3 homo s | 6 homo | 2 homo s | 4 homo | 7 homo s | 0 homo | 5 homo | B homo | 9 homo | 5 homo s | 6 homo s | 9 locust | 86 capri | 12084 cap: | 12080 capri | 12078 cap | 12088 cap | 12076 cap | 140,000 |

ALIGNMENTS

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Q93YC8;
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01-DEC-2001
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Q9QVE9;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UUN-2002 (TrEMBLrel. 21, Last annotation update)
protamine MP2 intermediate protein PMP2/11 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                   Chauviere M., Martinage A., Debarle M., Sautiere P., Chevaillier "Molecular characterization of six intermediate proteins in the processing of mouse protamine P2 precursor."; Eur. J. Biochem. 204:759-765(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10095;
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MEDLINE=92174934; PubMed=1541289;
 (TrEMBLrel. 19, (TrEMBLrel. 19, (TrEMBLrel. 24,
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                                                                                    PRELIMINARY;
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                Q9PRJ4;
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01-MAY-2000
01-MAY-2000
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01-MAR-2001 (Tr)
NAD+-dependent
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Q9H326;
Q1-MAR-2001
                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.

Kim Y.-O., Koh H.-J., Jo S.-H., Son M.-K., Huh T.-L.;

"Structural and functional characterization of the human NAD+-
dependent isocitrate dehydrogenase alpha subunit promoter.";

Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.

EMBL; AF157515; AAG43379.1; -.
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Nicotiana tabacum (Common tobacco).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
Smiids; Solanales; Solanaceae; Nicotiana.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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GO; GO:0019028; C:viral capsid; IEA.
GO; GO:0005198; F:structural molecul
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Matzke A.J.,
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ent isocitrate dehydrogenase 3 alpha subunit
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SEQUENCE
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Gene 235:131-137(1999).
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01-NOV-1999
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NCBI_TaxID=2885;
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SEQUENCE 9 AA; 1099 MW;
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MEDLINE=95380361; PubMed=7651903;
Conlon J.M., Platzack B., Marra L
"Isolation and biological activit
of the phylogenetically ancient f
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Amia calva (Bowfin).
Eukaryota; Metazoa; Chordata;
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raceae; Pylaiella.
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hone oxidoreductase subunit 3 (Fragment)
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GO; GO:0016563; F:transcriptional activator activity; IEA.

GO; GO:0045941; P:positive regulation of transcription; IEA

InterPro; IPR004247; Lentiviral Tat.

Pfam; PF02998; Lentiviral Tat; I.
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Turelli P., Guiguen F., Mornex J.-F., Vigne R.

"dUTPase minus CAEV is attenuated for pathoger
to A substitutions.";

Submitted (DEC-1996) to the EMBL/GenBank/DDBJ
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NCBI TaxID=11660;
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A Turelli P., Guiguen F., Mornex J.-F., Vigne R., Vullettelli P., GAEV is attenuated for pathogenesis and accumulate reductions.";

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RL Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.

EMBL; U81448; AAB60850.1; -.

DR GO; GO:0016563; F:transcriptional activator activity; IEA.

DR GO; GO:0045941; P:positive regulation of transcription; IEA.

DR GO; GO:0045941; P:positive regulation of transcription; IEA.

DR InterPro; IPR004247; Lentiviral Tat.

Pfam; PF02998; Lentiviral Tat; 1.
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"dUTPase minus CAEV is attenuated for pathogenesis and ac
to A substitutions.";
Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; U81432; AAB60818.1; -.
GO; GO:0016563; F:transcriptional activator activity; IEA
GO; GO:0045941; P:positive regulation of transcription; 1
InterPro; IPR004247; Lentiviral Tat.
Pfam; PF02998; Lentiviral Tat; I.
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Viruses; Retroid viruses; Retroviridae; Lentivirus
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SEQUENCE
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Turelli P., Guiguen F.,
"dUTPase minus CAEV is a
to A substitutions.";
                                                              Caprine arthritis encephalitis virus (CAEV).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI TaxID=11660;
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GO; GO:0016563; F:transcriptional activator activity; IEA.
GO; GO:0045941; P:positive regulation of transcription; IEA.
InterPro; IPR004247; Lentiviral Tat.
Pfam; PF02998; Lentiviral Tat; 1.
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Turelli P., Guiguen F., Morr
"dUTPase minus CAEV is atter
to A substitutions.";
Submitted (DEC-1996) to the
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Tat protein
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GO; GO:0016563; F:transcriptional
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GO; GO:0045941; P:positive regulation of transcription;
InterPro; IPR004247; Lentiviral Tat.
Pfam; PF02998; Lentiviral Tat; I.
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EMBL; U81446; AAB60846.1; -.
GO; GO:0016563; F:transcriptional activator activity; IEA.
GO; GO:0045941; P:positive regulation of transcription; IEA.
InterPro; IPR004247; Lentiviral Tat.
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Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
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Turelli P., Guiguen F., Mornex J.-F., Vigne R.

"dUTPase minus CAEV is attenuated for pathoger
to A substitutions.";

Submitted (DEC-1996) to the EMBL/GenBank/DDBJ
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Viruses; Retroid viruses; Retroviridae; Lentivirus
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SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

Turelli P., Guiguen F., Mornex J.-F., Vigne R., Querac J.-F.,

Turelli P., Guiguen F., Mornex J.-F., Vigne R., Querac J.-F.,

Turelli P., Guiguen F., Mornex J.-F., Vigne R., Querac J.-F.,

The All Submitted (DEC-1996) is attenuated for pathogenesis and accumulated accumulation of transcriptional accumulations. The EMBL; U81429; AAB60812.1; -J.

DR EMBL; U81429; AAB60812.1; -J.

DR GO; GO:0016563; F:transcriptional activator activity; IBA.

GO; GO:0045941; P:positive regulation of transcription; IEA.

DR InterPro; IPR004247; Lentiviral Tat.
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Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; U81447; AAB60848.1; -
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Viruses; Retroid viruses; Retroviridae;
NCBI_TaxID=11660;
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GO; GO:0016563; F:transcriptional activator activity; IEA.
GO; GO:0045941; P:positive regulation of transcription; IEA
InterPro; IPR004247; Lentiviral Tat.
Pfam; PF02998; Lentiviral Tat; 1.
NON TER 1 1 Turelli P., Guiguen F., Mornex J.-F., Vigne R., Querat G "dUTPase minus CAEV is attenuated for pathogenesis and a to A substitutions.";
Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases. 01-JUL-1997 01-JUL-1997 01-JUN-2003 SEQUENCE 012078 SEQUENCE FROM N.A. to A substitutions.";
Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; U81435; AAB60824.1; -. NCBI_TaxID=11660; Caprine arthritis Viruses; Retroid v SEQUENCE Turelli P., Guiguen "dUTPase minus CAEV Tat protein SEQUENCE FROM N.A. at protein CBI_TaxID=11660; Aprine arthritis encephalitis virus (CAEV) /iruses; Retroid viruses; Retroviridae; Len 19 9 œ 9 PGW 11 9 Similarity
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Last annotation update) 0 Score 24; Pred. No. Last sequence update)
Last annotation update) Created) 5A60BBB1E8644EB7 CRC64; 4B90BBB1E8644EB7 CRC64; Mismatches (CAEV) Vigne R., Querat pathogenesis and 12 12 DB 15; I . 2.2e+03; Lentivirus Lentivirus AA A Length 12; Length 12; Indels Indels G.; accumulates G.;
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                                                                                                                                                                                                         to A substitutions.";
Submitted (DEC-1996) to
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                                                                                                            SEQÜENCE
                                                                                                                         SEQUENCE FROM N.A. Turelli P., Guigue
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11660;
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Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; U81431; AAB60816.1; -.
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NCBI_TaxID=11660;
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"dUTPase minus CAEV
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100.0%; Pro
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01-NOV-1996 (TrEMBLrel. 0
01-JUN-2003 (TrEMBLrel. 2
Cytochrome b (Fragment).
Locusta migratoria (Migra
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EMBL; U81434; AAB60822.1; -.
GO; GO:0016563; F:transcriptional activator activity; IF
GO; GO:0045941; P:positive regulation of transcription;
InterPro; IPR004247; Lentiviral Tat.
Pfam; PF02998; Lentiviral Tat; I.
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Q34909;
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Viruses; Retroid v
NCBI_TaxID=11660;
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Submitted (DEC-1996) to the EMBL/GenBank/DDBJ
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01-JUN-2003
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SEQUENCE FROM N.A.

MEDLINE=88223478; PubMed=2836084;

McCracken A., Uhlenbusch I., Gellissen G.;

McCracken A., Uhlenbusch I., Gellissen G.;

"Structure of the cloned Locusta migratoria mitochondrial

restriction mapping and sequence of its ND-1(URF-1) gene."

Curr. Genet. 11:625-630(1987).

EMBL; X05286; CAA28905.1; -.

GO:0005739; C:mitochondrion; IEA.
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Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Acrididae; Oedipodinae; Locusta.
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01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
MHC class II antigen (Fragment).
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Q8MH35;
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Submitted (
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI_TaxID=9606;
                                                                                                               Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.; "HLA DQA1 gene generates multiple transcripts by alternative and polyadenylation of the 3' untranslated region."; Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.
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Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.;

"A new splicing acceptor site and polyadenylation sequence
contribute to increase the extraordinary diversity of DQA1
                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.,
Hoarau Splicing acceptor site and polyadenylation sequence
contribute to increase the extraordinary diversity of DQAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.; "HLA DQA1 gene generates multiple transcripts by alternative and polyadenylation of the 3' untranslated region."; Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.

EMBL, AF533934; AAM94848.1; -.
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Hoarau J.-J., Cesari M.,
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1; Mismatches
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Mammalia; Eutheria;
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Hoarau J.-J., Cesari M.,
"A new splicing acceptor
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Hoarau J.-J., Cesari M.,
"A new splicing acceptor
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A DQAl gene generates multiple transcripts by alternative
polyadenylation of the 3' untranslated region.";
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Eukaryota; Metazoa; Cl
Mammalia; Eutheria; Pı
NCBI_TaxID=9606;
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    EMBL; AFS
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SEQUENCE
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"A new splicing acceptor
contribute to increase th
isoforms.";
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                                                                                SEQUENCE FROM N.A.
Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.;
"HLA DQA1 gene generates multiple transcripts by alternative
and polyadenylation of the 3' untranslated region.";
Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.
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Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.,

"A new splicing acceptor site and polyadenylation sequence
contribute to increase the extraordinary diversity of DQA1
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Mammalia; Eutheria;
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Q8MH52;
Q8MH52;
01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation updat
                                                       EMBL; AFE
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HOARAU J.-J., Cesari M.M.,
"HLA-DQA1 genes generate mu
and polyadenylation of the
Submitted (MAY-2002) to the
                                                                                                                                         SEQUENCE FROM N.A.

Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.;

"HLA DQA1 gene generates multiple transcripts by alternative and polyadenylation of the 3' untranslated region.";

Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.
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"A new splicing acceptor
contribute to increase th
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1; Mismatches
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Pred. No.
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Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.,
"A new splicing acceptor site and polyadenylation sequence
contribute to increase the extraordinary diversity of DQA1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; (
Mammalia; Eutheria; E
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                                                                                                                   SEQUENCE
                                                                                                                                                                                                 Submitted
                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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                                                                                                                                             LA DQA1 gene generates multiple transcripts by alternative polyadenylation of the 3' untranslated region."; mitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.

L; AF533920; AAM94834.1; -.
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AF515810; AAM74917.1; -.
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3; Conservative
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Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.,
"A new splicing acceptor site and polyadenylation sequence contribute to increase the extraordinary diversity of DQA1 isoforms.";
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Hoarau J.-J., Cesari M.,
"A new splicing acceptor
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                                                                                                  ., Caillens H., Cadet F., Pabion M.; or site and polyadenylation sequence the extraordinary diversity of DQA1
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 Hoarau J.-J., Cesari M., Caillens H., Cadet F., I "HLA DQA1 gene generates multiple transcripts by and polyadenylation of the 3' untranslated region
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"HIA DQA1 gene generates multiple transcripts by alternative and polyadenylation of the 3' untranslated region."; Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; AF533902; AAM94816.1; -.
NON TER 1 1
SEQUENCE 14 AA; 1435 MW; C5EEAOFE15DF8EBD CRC64;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; !
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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MHC class II antigen (Fragment).
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Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.,
"A new splicing acceptor site and polyadenylation sequence
contribute to increase the extraordinary diversity of DQA1
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                                        SEQUENCE FROM N.A.

Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.;

"A new splicing acceptor site and polyadenylation sequence contribute to increase the extraordinary diversity of DQAI
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Mammalia; Eutheria;
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Hoarau J.-J., Cesari M.,
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Hoarau J.-J., Cesari M.,
"A new splicing acceptor"
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Mammalia; Eutheria;
NCBI_TaxID=9606;
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Hoarau J.-J., Cesari M., Caillens H., Cadet F., Pabion M.;

"HLA DQA1 gene generates multiple transcripts by alternative and polyadenylation of the 3' untranslated region.";

Submitted (JUL-2002) to the EMBL/GenBank/DDBJ databases.

EMBL; AF533910; AAM94824.1; -.

NON TER 1
SEQUENCE 14 AA; 1435 MW: CCDDACHONICE
                                                                                                                                                     EMBL;
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                                                                                                                                                                                                                                                        Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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01-OCT-2002 (TrEMBLrel. 22, Last
01-OCT-2002 (TrEMBLrel. 22, Last
MHC class II antigen (Fragment).
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CORZ_PERAM
ACHG_GRYBI
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P10939 nauphoeta c
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P14086 gryllus bim
P25418 libellula a
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P58805
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RESULT 2
CA32_LITCI STANDARD; PRT; 11 AA.

D CA32_LITCI STANDARD; PRT; 11 AA.

AC P82090;
AC P82090;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Caerulein 3.2/3.2Y4.
OS Litoria citropa (Australian blue mountains tree frog).
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| P38007 Chi | P42635 aed | P42634 aed | Ā08 80980d | P56264 lit | P82086 lit | P31351 vip | P31799 loc | | P30369 mac | | | |
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RESULT 1
CA31 LITCI STANDARD; PRT;
ID CA31 LITCI STANDARD; PRT;
AC P82089;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence up
DT 10-OCT-2003 (Rel. 42, Last annotation
DE Caerulein 3.1/3.1Y4.
                                                                                                                                                                                                                                                                                                                                                              WEDLINE-20057701; PubMed=10589099;

Wabnitz P.A., Bowie J.H., Tyler M.J.;

Wabnitz P.A., Bowie J.H., Tyler M.J.;

"Caerulein-like peptides from the skin glands of the Aus

"Caerulein-like peptides from the skin glands of the Aus

montains tree frog Litoria citropa. Part 1. Sequence det

using electrospray mass spectrometry.";

Rapid Commun. Mass Spectrom. 13:2498-2502(1999).

-!- FUNCTION: Hypotensive neuropeptide (Probable).

-!- SUBCELLULAR LOCATION: Secreted.

-!- TISSUE SPECIFICITY: Skin dorsal glands.

-!- PIM: Isoform 3.1Y4 differs from isoform 3.1 in not b
                                             Litoria citropa (Australian blue mountains tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hylidae;
Pelodryadinae; Litoria.
NCBI_TaxID=94770;
                                                                                                                                                                                                                                                                  -!- MASS SPECTROMETRY: MW=1407; METHOD=Electrospray.
-!- SIMILARITY: Belongs to the gastrin/cholecystokinin
InterPro; IPR001651; Gastrin.
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TISSUE=Skin secretion;
0; Mismatches
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Pred. No. 34;
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RESULT 3
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-!- MASS SPECTROMETRY: MW=1423; METHOD=Electrospray.
-!- SIMILARITY: Belongs to the gastrin/cholecystokinin family.
InterPro; IPR001651; Gastrin.
PROSITE; PS00259; GASTRIN; FALSE_NEG.
Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation;
                                                                                                                                                                                                                                                              Pandalus borealis (Northern red shrimp).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Caridea; Pandaloidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wabnitz P.A., Bowie J.H., Tyler M.J.; "Caerulein-like peptides from the skin glands of the Australian blue montains tree frog Litoria citropa. Part 1. Sequence determination
                                                                           PROSITE;
                                                                                                           -!- SUBCELLULAR LOCATION:
-!- SIMILARITY: Belongs to
                                                                                                                                                   "Structure of the red-pigment-concentrating Pandalus borealis.";
Biochim. Biophys. Acta 371:304-311(1974).
-!- FUNCTION: This hormone adapts the animal
                                                                                                                                                                                                                                              NCBI
                                                                                                                                                                                                                                                                                               01-NOV-1988 (Rel. 09, Created)
01-FEB-1994 (Rel. 28, Last sequence update
10-OCT-2003 (Rel. 42, Last annotation update
Red pigment concentrating hormone (RPCH)
                                                                                                                                                                                                                                                                                                                                            P08939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  using electrospray mass spectrometry.";
Rapid Commun. Mass Spectrom. 13:2498-2502(1999)
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Amphibia; Batrachia; An
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                         RPCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE, AN
                                                                                                                                                                                                  Fernlund P.;
                                                                                                                                                                                                            WEDLINE=75054965;
                                                                                                                                                                                                                        SEQUENCE.
                                                                                                                                                                                                                                                       Pandalidae;
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                                                                                     InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: Hypotensive neuropeptide (Probable).
SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Skin dorsal glands.
PTM: Isoform 3.2Y4 differs from isoform 3.2 in
                                                                                                                                           stimulating concentration
                                                                                               A61348; A61348.
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                                                                                                                                                                                                                                             TaxID=6703;
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Similarity
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                                                                           PS00256;
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1 AA;
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                                                                Amidation;
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                                                                            AKH;
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1363
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39.4%;
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                               AMIDATION.
86786775B9C44736 CRC64;
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Pred. No.
Score
Pred.
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                                                    Pyrrolidone PyrroLIDONE
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28;
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RESULT 4
HTF NAUCI
ID HTF NAUCI
AC P10939;
DT 01-ULL-198
DT 01-FEB-199
DT 10-OCT-200:
DE Hypertreha:
OS Leucophaea
OS Leucophaea
OS Leucophaea
OS Leucophaea
OS RIATTELLA
OC ENARYOTA;
OC Blaberidae
OC Neoptera;
OC Blaberidae
OC Neoptera;
OC SPECIES=N.
RY SEQUENCE.
RY Gaede G.,
RY Gaede G.,
RT Corpus Carn
RL Biochem. B
RN [2]
RP SEQUENCE.
RC SPECIES=L.
RC SPECIES=L.
RT Gromphador
RT B10. Chem
RN [2]
RT MEDLINE=91
RA WEDLINE=91
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                                                                                                                                Query Match
Best Local
                                                                                                       Matches
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SPECIES-L.maderae, G.portentosa, and B.germanica,

MEDLINE=90253659; PubMed=2340112;

Gaede G., Rinehart K.L. Jr.;

"Primary structures of hypertrehalosaemic neuropeptides isolated from the corpora cardiaca of the cockroaches Leucophaea maderae, the corpora cardiaca of Blattella germanica and Blatta orientalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Veenstra J.A., Camps F.;
"Structure of the hypertrehalosemic neuropeptide of the German cockroach, Blattella germanica.";
Neuropeptides 15:107-109(1990).
-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose major carbohydrate in the hemolymph of insects).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDITIVE SOLUTION (NEW YORK) Gaede G., Rinehart K.L. Jr.; Gaede G., Rinehart K.L. Jr.; Gaede G., Rinehart K.L. Jr.; "Amino acid sequence of a hypertrehalosaemic neuropeptide from the corpus cardiacum of the cockroach, Nauphoeta cinerea."; "Sinchem Biophys. Res. Commun. 141:774-781(1986).
                                                                                                                                                                                                                                                                                                                                                                                                PIR; A26381; A26381.
PIR; A60421; A60421.
PIR; S08997; S08997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gromphadorhina portentosa, Blattella germani
and of the stick insect Extatosoma tiaratum
atom bombardment mass spectrometry.";
Biol. Chem. Hoppe-Seyler 371:345-354(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-1989 (Rel. 11, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Hypertrehalosaemic hormone (HTH) (Hypertrehalosaemic neuropeptide).
Nauphoeta cinerea (Cinereous cockroach) (Gray cockroach),
Leucophaea maderae (Madeira cockroach),
Blattella germanica (German cockroach), and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SPECIES=B.germanica;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gromphadorina portentosa (Madagascan hissing cockro
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; i
                                                                                                                                                                                                                                                                                           Neuropeptide; Amidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=91179584; PubMed=2080017;
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                                                                                                                                                                                                                                                                                                                       InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
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                                                                                                                                                                                                                                                                                                                                                                            S08998; S08998.
3 NFSPGW
                                                  6 HYGPGW 11
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3; Conserv
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                                                                                                                                                                                                               10 AA;
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6990, 6988, 6973,
                                                                                                       Conservative
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                                                                                                                                                                                                               1092 MW;
                                                                                                                                39.4%;
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                                                                                                                                  Score 28;
Pred. No.
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                                                                                                       <u>ب</u>
                                                                                                                                                                                                               CRC64;
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Blaberoidea;
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GON2_CHICK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Distanded dogfish bit Natl. /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Squalus acanthias (Spiny dogfish),
Hydrolagus colliei (Spotted ratfish)
Clupea pallasii (Pacific herring)
Eukaryota; Metazoa; Chordata; Craniat
                                                                                                                                                     SEQUENCE, AND FUNCTION.

SPECIES-C.pallasii; TISSUE-Brain, and Pituitary;

MEDLINE=20114351; PubMed=10650929;

Carolsfeld J., Powell J.F.F., Park M., Fischer W.H., Craig A.G.,

Chang J.P., Rivier J.E., Sherwood N.M.;

"Primary structure and function of three gonadotropin-releasing hormones, including a novel form, from an ancient teleost, herri Endocrinology 141:505-512(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alligator mississippiensis (American
                                                                                                                                                                                                                                                                                                                                                                                                                          of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=91352338; PubMed=1882082;
Lovejoy D.A., Fischer W.H., Parker D.B., McRory J.E., Park M.,
Lance V., Swanson P., Rivier J.E., Sherwood N.M.;
"Primary structure of two forms of gonadotropin-releasing hormone from brains of the American alligator (Alligator mississippiensis)
Regul. Pept. 33:105-116(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
-!- FUNCTION: Stimulates the secretion of g-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the GnRH family.
PIR; A61126; A61126.
PIR; B46030; B46030.
PIR; B60066; RHAQ2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SPECIES=H.colliei; TISSUE=Brain; MEDLINE=91340067; PubMed=1678723; Lovejoy D.A., Sherwood N.M., Fisc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lovejoy D.A., Fischer W.H., Ngamvongchon S., Craig A.G. Nahorniak C.S., Peter R.E., Rivier J.E., Sherwood N.M.; "Distinct sequence of gonadotropin-releasing hormone (G dogfish brain provides insight into GnRH evolution."; proc. Natl. Acad. Sci. U.S.A. 89:6373-6377(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probably controlled by two in avian species.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  chicken hypothalamus:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=84222059;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Primary structure of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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Aves; Neognathae; Galliformes;
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; PubMed=6427779;
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                                                                                                                                                                                                                                                                                                                                                                                              of gonadotropin-releasing hormone (ratfish: Hydrolagus colliei)."; nol. 82:152-161(1991).
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gonadotropin-releasing
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P25423;

P25423;

01-MAY-1992 (Rel. 22, Created)

01-FEB-1994 (Rel. 28, Last sequence upda

01-FEB-1994 (Rel. 42, Last annotation up

10-OCT-2003 (Rel. 42, Last annotation up

Adipokinetic hormone (AKH).

Melolontha melolontha (Cockchafer),

Melolontha melolontha (Fockchafer),

Geotrupes stercorosus (Dor beetle), and

Geotrupes stercorosus (Flower beetle).

Pachnoda marginata (Flower beetle).

Eukaryota; Metazoa; Arthropoda; Hexapod

Cockchafera; Fndopterygota; Coleoptera; Po

and

poda; Insecta; Pterygota; Polyphaga; Scarabaeiformia;

STANDARD;

PRT;

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on update)

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RESULT
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AKH_MEI
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01-MAY-1992
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MOD_RES
MOD_RES
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P27429;
01-AUG-1992
                                                                                                                                                                                                                                                                                                                                            MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lovejoy D.A., Fischer W.H., Ngamvongchon S., Craig A.G., Nahorniak C.S., Peter R.E., Rivier J.E., Sherwood N.M.; "Distinct sequence of gonadotropin-releasing hormone (Gn dogfish brain provides insight into GnRH evolution."; Proc. Natl. Acad. Sci. U.S.A. 89:6373-6377(1992).

-!- FUNCTION: Stimulates the secretion of gonadotropins.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the GnRH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SQUAC
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28-FEB-2003 (Rel. 41, Last annotation updat
Gonadoliberin (Gonadotropin-releasing horm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Squalea; Squaloidei; Squalidae; Squalus.
                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIR; A46030; A46030.
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                                                                                                                                                                                                                                                                                                                                                                                                                                interPro; IPRUUZULZ;
Pfam; PF00446; GnRH;
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PS00473; GNRH; 1.
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Best Local S
Matches 3
 MEDLINE=90046758; PubMed=2813385;

Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachma Vogel V.W., Zhang Y.-S., Hayes D.K.;

"Primary structure of two neuropeptide hormones with hypotrehalosemic activity isolated from the corpora cflies (Diptera).";

Proc. Natl. Acad. Sci. U.S.A. 86:8161-8164(1989).

-!- FUNCTION: This hormone, released from cells in the cardiaca after the beginning of flight, causes rediglycerides from the fat body and then stimulate
                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JAN-1990
01-FEB-1994
10-OCT-2003
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Spectrometry.";

Biol. Chem. Hoppe-Seyler 373:133-142(1992).

-!- FUNCTION: This hormone, released from ce cardiaca after the beginning of flight, diglycerides from the fat body and then diglycerides as an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuropeptide;
MOD_RES
MOD_RES
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"Primary structures of neuropeptides isolated from the corpora cardiaca of various cetonid beetle species determined by pulsed-liquid phase sequencing and tandem fast atom bombardment mass
                                                                                                                                                                                                 TISSUE=Corpora cardiaca;
MEDLINE=90046758; PubMed
                                                                                                                                                                                                                                                                                                                   (DCC 1).
Tabanus atratus (Horse fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Metazoa; Arthropoda; Brachycera; Tabanomorpha; "
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                                                                                                                                                                                                                                                                                   NCBI_TaxID=7207;
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S15422; S15422.
S21663; S21663.
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2265187; PubMed=1586453;
Lopata A., Kellner R., Rir
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hormone (AKH)
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the AKH / HRTH / RPCH
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1; Mismatches
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867AB775AB544736 CRC64;
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corpora
ease of
the flight
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GON3_ONCKE
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Best Local
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P20367; P817
01-FEB-1991
                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES-C.pallasii; TISSUE-Brain, and Pituitary; MEDLINE-20114351; PubMed-10650929; Carolsfeld J., Powell J.F.F., Park M., Fischer W.H., Craig A.G., Chang J.P., Rivier J.E., Sherwood N.M.; "Primary structure and function of three gonadotropin-releasing hormones, including a novel form; from an ancient teleost, herr: Endocrinology 141:505-512(2000).
                                                                                                                                                                                    MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Clupea pallasii (Pacific herring).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                 PIR;
                                                                                                                                                                                                                                                                                                                               -!- SUBCELLULAR LOCATION:
-!- SIMILARITY: Belongs to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sherwood N., Eiden L., Brownstein M., "Characterization of a teleost gonadot Proc. Natl. Acad. Sci. U.S.A. 80:2794-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SPECIES=O.keta;
MEDLINE=83195140;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-FEB-1991 (Rel. 17, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Gonadoliberin III (Gonadotropin-releasing hormone
RH III) (Luliberin III).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            muscles to use these diglycerides as an energy source.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
PIR; A33995; A33995.
InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation; Flight; Pyrrolidone carboxylic ac
                                                                                                                                                                                                                                                 PROSITE;
                                                                                                                                                                                                                                                                                           InterPro;
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3; Conserv
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3; Conserv
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                                                                                                                                                                                                                         Amidation; Hypothalamus;
QHWSYGW
                                                                                                                                                                                                                                                00446; GnRH; 1.
PS00473; GNRH; 1.
                                       KHYGPGW
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42.9%;
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                                                                             Score 25; DB Pred. No. 1.6e 2; Mismatches
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Pred. No.
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                                                                                                                                                                                     AMIDATION
                                                                                                                                                                                                      mus; Pyrrolidone carboxylic PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tein M., Spiess J., Rivier J., Vale t gonadotropin-releasing hormone."; 80:2794-2798(1983).
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                                                                        JB 1.6e+02;
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follicle-stimulating
                                                                                                                                                                 CRC64;
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Best Local S
Matches 3
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Q9PRZ1;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last seque
16-OCT-2001 (Rel. 40, Last annot
Lysyl-bradykinin-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ONCMY
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Eukaryota; Metazoa; Arthropoda; He
Neoptera; Endopterygota; Diptera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P14596;
01-JAN-1990 (Rel. 13,
01-FEB-1994 (Rel. 28,
10-OCT-2003 (Rel. 42,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=90046758; PubMed=2813385; Jaffe H., Raina A.K., Riley C.T., Vogel V.W., Zhang Y.-S., Hayes D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypertrehalosaemic
(DCC II).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Proc. Natl. Acad. Sci. U.S.A.
trout plasma.";
FEBS Lett. 334:75-78(1993).
-!- FUNCTION: SMOOTH MUSCLE CONTRACTION. PROBABLY PLAYS
THIS SYSTEM IN CARDIOVASCULAR REGULATION IN FISH.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Plasma.
-!- SIMILARITY: Belongs to the bradykinin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Neuropeptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Primary structure of two neuropeptide hormones with adipokinetic and hypotrehalosemic activity isolated from the corpora cardiaca of horse flies (Diptera).";
                                                                                                                                                              MEDLINE=94039817; PubMed=8224232; Conlon J.M., Olson K.R.; "Purification of a vasoactive per
                                                                                                                                                                                                                                                                                                   Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EQUENCE.
CISSUE=Corpora
                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             major carbohydrate in the hemolymph of insects).
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the AKH / HRTH / RPCH family.; B33995; B33995.; B33995.
erPro; IPR002047; AKH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . Natl. Acad. Sci. U.S.A. 86:8161-8164(1989). FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7 YGPGW 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
3; Conserv
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yota; Diptera; Brachycera; Tabanomorpha;
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Last annotation update)
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916036786771A9D1 CRC64;
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Bradykinin; Vasodilator
SEQUENCE 10 AA; 1193
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STRAIN=Wistar; TISSUE=Heart, and Liver;

STRAIN=95324529; PubMed=7601105;

MEDLINE=95324529; PubMed=7601105;

Schaegger H., Noack H., Halangk W., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Halangk T., Brandt U., von Jagow Schaegger H., Noack H., Halangk T., Ha
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P80432;
01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence upon 1-NOV-1995 (Rel. 41, Last annotation upon 1-NOV-19
                                                                                                                                    Thunnus obesus (Bigeye tuna).
Eukaryota; Metazoa; Chordata; Craniata;
Actinopterygii; Neopterygii; Teleostei;
Acanthomorpha; Acanthopterygii; Percomor
Scombridae; Thunnus.
NCBI TaxID=8241;
                                                                                                                                                                                                                                                                                                              P80982;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence up
16-OCT-2001 (Rel. 40, Last annotation
Cytochrome c oxidase polypeptide VIIc
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PIR;
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-!- FUNCTION: This protein is one of the rechains of cytochrome c oxidase, the temitochondrial electron transport.

-!- CATALYTIC ACTIVITY: 4 ferrocytochrome
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Mammalia; Eutheria;
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COX7C OR COX7C1.
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SIMILARITY: Belongs to
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TISSUE=Heart,

and Liver;

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Matches 5
                                                                                                                        SEQUENCE.
SPECIES=L.decemlineata; TISSUE=Corpora
SPECIES=1.decemlineata; TISSUE=Corpora
MEDLINE=90160053; PubMed=2576128;
Gaede G., Kellner R.;
Gaede G., Kellner R.;
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P04548;
13-AUG-1987
01-FEB-1994
10-OCT-2003
   SPECIES=B.orientalis;
                                                                                                                                                                                                                                                            "Isolation and primary structure of two peptides with cardioacceleratory and hyperglycemic activity from th cardiaca of Periplaneta americana.";
Proc. Natl. Acad. Sci. U.S.A. 81:5575-5579(1984).
                             SEQUENCE
                                                                                                beetle and
                                                                                                                                                                                                                                                                                                                                                         MEDLINE=84298179; PubMed=6591205; Scarborough R.M., Jamieson G.C., Miller C.A., Schooley D.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Structures of two cockroach neuropeptides assigned bombardment mass spectrometry."; Biochem. Biophys. Res. Commun. 124;350-358(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES=P.americana;
MEDLINE=85046530; PubMed=6548628;
Witten J.L., Schaffer M.H., O'Shea M.,
Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-1987 (Rel. 05, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
10-OCT-2003 (Rel. 42, Last annotation update)
Hypertrehalosaemic factor I (Neuropeptide M-I) (Periplanetin CC-I)
(PeA-CAH-I) (LeD-CC-I) (Hypertrehalosaemic neuropeptide I).
Periplaneta americana (American cockroach),
Leptinotarsa decemlineata (Colorado potato beetle), and
                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES=P.americana;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eur.
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                                                                       Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blatta orientalis (Oriental cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PERAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Blattidae; Periplaneta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oxidoreductase; Inner membrane; Mitochondrion NON_TER 10 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=97454291; PubMed=9310366; Arnold S., Lee I., Kim M., Song E.,
                                                                                                                    The metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Neoptera; Orthopteroidea; Dictyoptera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The subunit structure of cytochrome-c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         c + 2 H(2)O.
SUBCELLULAR LOCATION: Mitochondrial
SIMILARITY: Belongs to the cytochron; S77990; S77990.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: This protein is one of the nuclear-coded polypeptide chains of cytochrome c oxidase, the terminal oxidase in mitochondrial electron transport.

CATALYTIC ACTIVITY: 4 ferrocytochrome c + O(2) = 4 ferricytochrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 S.,
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                                                                     abolic neuropeptides of the nd the American cockroach as 10:1287-1289(1989).
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71.4%;
TISSUE=Corpora cardiaca;
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Pred. No.
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                                                                                            corpus cardiacum from the re identical.";
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                                                                                                                                                                                           cardiaca;
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5.5e+02;
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Blattoidea;
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                                                                                                                    potato
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RESULT
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Best Local S
Matches 2
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P25419;
01-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VI-MAY-1992 (Rel. 22, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Hypertrehalosaemic factor (HOTH) (Hypertrehalosemic neuropeptide).
Tenebrio molitor (Yellow mealworm), and
Zophobas rugipes.
                                                                                                                      Neuropeptide;
MOD_RES
MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=90253659; PubMed=2340112;
Gaede G., Rinehart K.L. Jr.;
Primary structures of hypertrehalosaemic neuropeptides isolated from the corpora cardiaca of the cockroaches Leucophaea maderae,
Gromphadorhina portentosa, Blatta orientalis and of the stick insect Extatosoma tiaratum assigned by tandem fast and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry.";
Biol. Chem. Hoppe-Seyler 371:345-354(1990).
-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose is the major carbohydrate in the hemolymph of insects).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
                                                                                                                                                                                                                                                                       PIR;
                                                                                                                                                                                                                                                                                                                                                                                          "The primary structure of the hypertrehalosemic neuropeptide from tenebrionid beetles: a novel member of the AKH/RPCH family."; Peptides 11:455-459(1990).
                                                                                                     SEQUENCE
                                                                                                                                                                                                InterPro; IPR002047; AKH. PROSITE; PS00256; AKH; 1.
                                                                                                                                                                                                                                                 PIR;
                                                                                                                                                                                                                                                                                         elevate the level of trehalose in the hemolymph (tre
major carbohydrate in the hemolymph of insects).
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AXH / HRTH / RPCH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SPECIES=T.molitor, and Z.rugipes; TISSUE=Corpora cardiaca; MEDLINE=90341081; PubMed=2381871;
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Tenebrionidae; Tenebrio.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaede G., Rosinski G.;
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, A44960, A44960.
, A49823, A49823.
, S08995, S08995.
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B43976;
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     Similarity 33. 2; Conservative
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2; Conserv
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                                                                                                     α
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Endopterygota; Coleoptera; Polyphaga; Cucujiformia;
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                                                                                                   AA,
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                                                                                                                                                                                                                                                                                                                                           Hypertrehalosaemic factors are neuropeptides he level of trehalose in the hemolymph (trehalohydrate in the hemolymph of insects).
                                                                                                                                                                        Amidation;
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1005 MW;
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                         31.0%;
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                                                                                                                                                Pyrrolidone carboxylic acid.
PYRROLIDONE CARBOXYLIC !
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     Score 22; DB Pred. No. 1.4e 2; Mismatches
                                                                                                   AMIDATION.
86745775B9C44736 CRC64;
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.4e+05;
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CORZ_PERAM
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N1_PETMA
GON1_F
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20-MAR-1987 (Rel. 04, Last sequence updated to the sequence of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P11496;
01-OCT-1989
01-FEB-1994
10-OCT-2003
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MOD_RES
MOD_RES
SEQUENCE
                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Arthropoda;
Neoptera; Orthopteroidea; Dictyo;
Blattidae; Periplaneta.
NCBI_TaxID=6978;
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Eukaryota; Metazoa; Chordata; Crania:
Petromyzontiformes; Petromyzontidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P04378;
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Sherwood N.M., Sower S.A., Marshak D.R., Fraser B.A.,
"Primary structure of gonadotropin-releasing hormone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=7757;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        brain.";
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"Isolation and structure of corazonin, the American cockroach."; FEBS Lett. 250:231-234(1989).
-i- FUNCTION: Cardioactive peptide. Coin the physiological regulation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIR;
                                                                                                                                                                                 TISSUE=Corpora cardiaca;
MEDLINE=89325572; PubMed=2753132;
                                                                                                                                                                                                                                                   SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIR; A01412; RHLMGS.
InterPro; IPR002012; GnRH.
Pfam; PF00446; GnRH; 1.
PROSITE; PS00473; GNRH; 1.
Hypothalamus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Corazonin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the GRR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FUNCTION:
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3; Conserv
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10
10 AA;
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(Rel. 28, Last seq
(Rel. 42, Last ann
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                261:4812-4819(1986).
Stimulates the secretion
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Chordata; Craniata; Vertebrata;
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42.9%;
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                                                                                                                                                                                                                                                                                                                                                                     erican cockroach).
opoda; Hexapoda; Insecta;
Dictyoptera; Blattaria; !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence update)
annotation update)
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1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         imus; Pyrrolidone carboxylic acid.
PYRROLIDONE CARBOXYLIC ACID.
AMIDATION.
1E4B36237B1735AB CRC64;
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lon update)
ssing hormone
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       Corazonin is of the heart
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5.2e+02;
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follicle-stimulating
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01-JAN-1990 (Rel. 13, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation updat
Adipokinetic hormone G (AKH-G) (RO II).
Gryllus bimaculatus (Two-spotted cricket),

update)

STANDARD;

Contact Metazou, Eukaryota; Metazou, Eukaryota; Metazou, Conthopteroidea;

Romalea microptera (Lubber grasshopper). Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Orthoptera; Ensifera; Gryllidae; Gryllinae;

(Two-spotted cricket), and

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RESULT 19
AKHG GRYBI
ID AKHG GRYBI
AC P14086;
DT 01-JAN-1990 (
DT 01-FEB-1994 (
DT 10-OCT-2003 (
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BOML_PSEGU
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Best Local (
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PIR; S05002; S05002.

Neuropeptide; Amidation; Pyrrolidon
MOD_RES 1 1 1 PYRROL
MOD_RES 11 11 AMIDAT
SEQUENCE 11 AA; 1387 MW; C7CFF3
                                                                                                                                                                                                                                                                                                                                                                                                         MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=90287814; PubMed=2356157;
Simmaco M., Severini C., de Biase D., Barra |
Roberts J.D., Melchiorri P., Erspamer V.;
"Six novel tachykinin- and bombesin-related |
the Australian frog Pseudophryne guntheri.";
Peptides 11:299-304(1990).
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pseudophryne guentheri (Guenther's toadlet).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
Myobatrachinae; Pseudophryne.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
Bombesin-like peptide 1 (PG-L)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR000874; Bombesin.
Pfam; PF02044; Bombesin; 1.
PROSITE; PS00257; BOMBESIN; 1.
Amphibian defense peptide; Bombesin
Pyrrolidone carboxylic acid.
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-!- SIMILARITY: Belongs to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE.
TISSUE=Skin secretion;
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                                                                                                                                                                                                                                                                                                                                                                                                             13 AA;
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1372 MW;
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75.0%;
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PYRROLIDONE CARBOXYLIC ACID
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Pred. No.
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Pred. No. 6.8e+02;
0; Mismatches 1
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C7CFF32D6415AB46 CRC64;
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D6DE0D24BD98C366
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5.7e+02;
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                                                                                                    AA.
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Myobatrachidae;
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Best Local S
Matches 2
                                                                                                                                                                                                                                                                                                                                                                                                   rue putative ancestral peptide of the adipokinetic/red-pigment-concentrating hormone family isolated and sequenced from a dragonfly.";
Biol. Chem. Hoppe-Savian and Sequenced from a sequenced fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AKH_LIBAU
P25418;
01-MAY-1992
                                                                                                                    -!- FUNCTION: This hormone, released from cells in the corpora cardiaca after the beginning of flight, causes release of diglycerides from the fat body and then stimulates the flig muscles to use these diglycerides as an energy source.
-!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
PIR; S10596; S10596.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SPECIES=R.microptera; TISSUE=Corpora cardiaca;
MEDLINE=89145002; PubMed=3226948;
Gaede G., Hilbich C., Beyreuther K., Rinehart K.L. Jr.;
"Sequence analyses of two neuropeptides of the AKH/RPCH-family
the lubber grasshopper, Romalea microptera.";
Peptides 9:681-688(1988).
-!- FUNCTION: This hormone, released from cells in the corpora
cardiaca after the beginning of flight, causes release of
diglycerides from the fat body and then stimulates the flig
muscles to use these diglycerides as an energy source.
-!- SUBCELIULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.

DIE. A28004: A28004
                                           InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-1992 (Rel. 22, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Adipokinetic hormone (AKH).
Libellula auripennis (Skimmer dragonfly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Palaeoptera; Odonata; Anisoptera; Libellulidae; Libellula.
Neuropeptide; Amidation; Flight;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=90359055;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Corpora cardiaca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE,
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MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID.

MOD_RES 8 AMIDATION.

SEQUENCE 8 AA; 938 MW; 867861B5B9C452D6 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=6966;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gryllus.
NCBI_TaxID=6999,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=2390213;
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Pred. No.
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   Pyrrolidone carboxylic
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1.4e+05;
2;
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Matches 2
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Best Local
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                                                                                                                             Canis familiaris (Dog).
Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Carnivora; Fissipedia; Canidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Corpora cardiaca;
MEDLINE=89145002; PubMed=3226948;
Gaede G., Hilbich C., Beyreuther K., Rinehart K.L. Jr.;
Gaede G., Hilbich C., Beyreuther K., Rinehart K.L. Jr.;
"Sequence analyses of two neuropeptides of the AKH/RPCH-family from the lubber grasshopper, Romalea microptera.";
Peptides 9:681-688(1988).
-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose is the content of the level of trehalose in the hemolymph (trehalose is the content of the level of trehalose in the hemolymph (trehalose is the content of the level of trehalose in the hemolymph (trehalose is the content of the level of trehalose is the level of the level of trehalose is the content of the level of the level of trehalose is the level of the level 
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ROMMI
HTF1 RO
P18110;
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MOD_RES
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TISSUE=Heart;
MEDLINE=98163340;
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10-OCT-2003
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01-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha; Acridoidea; Romaleidae; Romalea.
                                                                                                                   NCBI_TaxID=9615
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PROSITE; PS00256; AKH; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          major carbohydrate in the hemolymph of insects).
SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
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RB-1994 (Rel. 28, Last sequence up
R-2003 (Rel. 42, Last annotation
(Hypertrehalosaemic factor).
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2; Conserv
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                                                                                                                                                                                                                                                                        (Rel. 34, Created)
(Rel. 34, Last sequence update)
(Rel. 42, Last annotation update)
none oxidoreductase 42 kDa subunit (EC 1)
(Complex I-42KD) (CI-42KD) (Fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
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PubMed=9504812;
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2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8665A771A9C452D6 CRC64;
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Pred. No. 1.4e+05;
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                                                                                                                                                    Euteleostomi, Canis.
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RESULT 24
HTF2_PERAM
ID HTF2_PERAM
AC P04549;
DT 13-AUG-1987
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RF1_CONSP
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Matches 4
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28-FEB-2003
28-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dog heart proteins.";
Electrophoresis 18:2795-2802(1997).
-!- FUNCTION: Transfer of electrons from NADH to the chain. The immediate electron acceptor for the
                                                                                                                                                                  Neurotoxin; Toxin;
MOD_RES 12
SEQUENCE 12 AA;
                                                                                                                                                                                                                                                                             "Conorfamide, a Conus venom peptide belonging neuropeptides.";
                                                                                                                                                                                                                                                                                                    MEDLINE=21605839; PubMed=11738233;
Maillo M., Aguilar M.B., Lopez-Vera E.,
Olivera B.M., Heimer de la Cotera E.P.;
                                                                                                                                                                                                                                                                                                                                                                 Apogastropoda; Caenogastropoda; Gastropoda; Orthogastropoda;
Neogastropoda; Caenogastropoda; Sorbeoconcha; Hypsogastropo
Neogastropoda; Conoidea; Conidae; Conus.
NCBI_TaxID=192919;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P58805;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oxidoreductase; NAD; Ubiquinone; Flavoprotein; FAD; Mitochondrion. NON_TER 12 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dunn M.J., C
                                                                                                                                                                                                                                                                                                                                                                                                                                Conorfamide-Sr1
                                                                                                                                                                                                                                                                                                                                                  SEQUENCE,
                                                                                                                                                                                                                                                                                                                                         SSUE=Venom;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chain. The immediate electron acceptor for the to be ubiquinone.

CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+)
CATALYTIC ACTIVITY: NADH + acceptor = NAD(+) +
COFACTOR: Binds 1 FAD per subunit.

SUBUNIT: Mammalian complex I is composed of 45
This a component of the hydrophobic protein fra
SUBCELLULAR LOCATION: Mitochondrial matrix.
                                                                                                                                                                                                                        FUNCTION: Causes hyperactivity in mice.
SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expressed by the venom duct.
MASS SPECTROMETRY: MW=1454.8; METHOD=Electrospray
                                                                                                                                                                                                              SIMILARITY: Belongs to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CONSP
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                                                                                                                      Similarity 4; Conserv
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                                                                            GPMGW
                                                                                                   GP-GW
                                                                                                                                                                                                                                                                                                                                                 SYNTHESIS,
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                                                                                                                                                                                                                                                                                                                                                                                                                                      (Rel. 41, Created)
(Rel. 41, Last seq
(Rel. 41, Last ann
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100.0%;
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  Created)
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                                                                                                                        Score 19.5; I
Pred. No. 1.7e
0; Mismatches
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Pred. No.
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                                                                                                                                                                   AMIDATION.
2510671E49D772D3
                                                                                                                                                                                                                                                                                                                                                  SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT:
                                                                                                                                                                                                              FARP (FMRFamide related peptide)
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on update)
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1.4e+0
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                                                                                                                                 .7e+03;
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AD(+) + reduced acce
                                                                                                                                            A.G.,
                                                                                                                                                                                                                                                                                                                                                                                              Hypsogastropoda;
                                                                                                                                                                     CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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                                                                                                                                            Length 12;
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nzyme is believed
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A Gaede G., Rinehart K.L. Jr.;
The corpora cardiaca of the cockroaches Leucophaea maderae,
The corporation of the stick in the cockroaches Leucophaea maderae,
The corporation cardiaca of the cockroac
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Best Local S
Matches 2
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Hypertrehalosaemic factor II (Neuropeptide M-II) (Periplanetin (PeA-CAH-II) (LeD-CC-II) (Hypertrehalosaemic neuropeptide II).
Periplaneta americana (American cockroach),
Leptinotarsa decemlineata (Colorado potato beetle), and
Blatta orientalis (Orientalia cockroach).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SPECIES=P.americana;
MEDLINE=85046530; PubMed=6548628;
Witten J.L., Schaffer M.H., O'Shea M.,
Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                           -!- SUBCELLULAR LOCATION: S

-!- SIMILARITY: Belongs to

PIR; B44960; B44960.

PIR; B49823; B49823.

PIR; S08996; S08996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SPECIES=L.decemlineata; TISSUE=Corpora MEDLINE=90160053; PubMed=2576128; Gaede G., Kellner R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Isolation and primary structure of two peptides cardioacceleratory and hyperglycemic activity frocardiaca of Periplaneta americana.";
Proc. Natl. Acad. Sci. U.S.A. 81:5575-5579(1984)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=84298179; PubMed=6591
Scarborough R.M., Jamieson G.
Miller C.A., Schooley D.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; F
Blattidae; Periplaneta.
NCBI TaxID=6978, 7539, 6976;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES=B.orientalis; TISSUE=Corpora MEDLINE=90253659; PubMed=2340112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The metabolic neuropeptides of the beetle and the American cockroach as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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10-OCT-2003 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE.
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 FTPNW
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ilarity 40.0%;
Conservative
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Jamieson G.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28, Last sequence update)
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                                                                            Score 19; DB
Pred. No. 1.4e
1; Mismatches
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                                                                                                                                                                               AMIDATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ne corpus cardiacum are identical.";
                                                                                                                                                                                                  DONE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cardiaca;
                                                                        1.4e+05;
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                                                                             Indels
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RESULT 25
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MOD_RES 1

MOD_RES 10 1

SEQUENCE 10 AA, 12
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28-FEB-2003 (Rel
Gonadoliberin II
(Luliberin III).
                                                                  Gaede G., Rinehart K.L. Jr.; "Primary structure of the hypertrehalosaemic factor II for the corpus cardiacum of the Indian stick insect, Carausius modetermined by fast atom bombardment mass spectrometry."; bicl. Chem. Hoppe-Seyler 368:67-75(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sower S.A., Chiang Y.-C., Lovas S., Conlon "Primary structure and biological activity releasing hormone from lamprey brain."; Endocrinology 132:1125-1131(1993).
           SPECIES=E.tiaratum; TISSUE=Corpora MEDLINE=90253659; PubMed=2340112; Gaede G., Rinehart K.L. Jr.;
                                                                                                                         SPECIES=C.morosus; TISSUE=Corpora
MEDLINE=87157103; PubMed=3828078;
                                                                                                                                                                     Neoptera; Orthopteroidea;
Heteronemiidae; Carausius.
NCBI TaxID=7022, 7024;
                                                                                                                                                                                             Carausius morosus (Indian stick insect), and Extatosoma tiaratum (Stick insect).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Neoptera; Orthopteroidea; Phasmatodea; Euphasmida;
                                                                                                                                                                                                                                           Hypertrehalosaemic factor neuropeptide II).
                                                                                                                                                                                                                                                                  01-JUL-1989 (Rel. 11, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation updat
                                                                                                                                                                                                                                                                                                                  HTF2
                                                                                                                                                                                                                                                                                                                            CARMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Brain;
MEDLINE=93178316; PubMed=8440174;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Hyperoartia;
Petromyzontiformes; Petromyzontidae; Petromyzon.
                                             SEQUENCE.
                                                                                                                                                    SEQUENCE.
                                                                                                                                                                                                                                                                                          P11385;
01-JUL-1989
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01-JUL-1993
"Primary
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                                                                                                                                                                                                                                                                                                                 CARMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PF00446; GnRH;
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                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 2; Conserv
  structures
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(Rel. 41, Last annotation updat)
in_III (Gonadotropin-releasing)
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2012; GnRH.
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28.6%;
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hypertrehalosaemic neuropeptides
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Pred. No. 1.7e
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284B36237AA1F5A3
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                                                                                                                                        cardiaca;
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(HRTH-II)
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1.7e+03;
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follicle-stimulating
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                                                                                        factor II from the Carausius morosus,
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e a third
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CRC64;
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                                                                                                                                                                                                                                                       (Hypertrehalosaemic
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Phasmatoidea;
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P SEQUENCE:
C TISSUE=Corpora cardiaca;
CX MEDLINE=88326324; PubMed=3415690;
UX MEDLINE=88326324; PubMed=3501690;
UX MEDLINE=88326324; PubMed=3501690;
UX MEDLINE=88326324; PubMed=3501690;
UX MEDLINE=88326324; PubMed=3501690;
UX MEDLIN
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HTF_HELZE
ID HTF_HELZE
AC P16353;
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Best Local
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MEDLINE=93129188; PubMed=1482345;
Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;
"A tryptophan-substituted member of the AKH/RPCH family isolated a stick insect corpus cardiacum.";
Biochem. Biophys. Res. Commun. 189:1303-1309(1992).

-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose imajor carbohydrate in the hemolymph of insects).

-!- SUBCELIULAR LOCATION: Secreted.

-!- SUBCELIULAR LOCATION: Secreted.

-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
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MOD RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hypertrehalosaemic hormone (HeZ-HRTH).
Heliothis zea (Corn earworm) (Bollworm).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia; Noc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the corpora cardiaca of the cockroaches Leucophaea maderae, Gromphadorhina portentosa, Blattella germanica and Blatta orientalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry."; along the characteristic constants of the complex of the stick insect of the stick insect of the stick insect of the constant of the stick insect of the corporate of the corporate
                                      PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation;
MOD_RES 1 1
MOD_RES 10 10
SEQUENCE
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01-FEB-1994 (Rel. 28, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
Hypertrehalosaemic hormone (HeZ-HRTH).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Neuropeptide; Amidation; MOD_RES 1 1
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SPECIES=C.morosus; TISSUE=Corpora cardiaca;
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interPro; 1:...
ps00256;
                                                                                                                                                                InterPro; IPR002047; AKH. PROSITE; PS00256; AKH; 1.
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S09138; S09138
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2; Conserv
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1096
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Pred. No.
                                                                         PYRROLIDONE CARBOXYLIC
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AMIDATION
8E70367865A5B9D1 CRC64;
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STRAIN=168 / JH642;
STRAIN=168 / JH642;
Graumann P.L., Schmid R
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_CSI5_I
SEQUENCE.

SPECIES=Horse;

Dixon J.S., Li

"The isolation
                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=168 / JH642;
MEDLINE=96345629; PubMed=8755892;
Graumann P., Schroeder K., Schmid R., Marahiel
"Cold shock stress-induced proteins in Bacillu
"Cold shock stress-induced proteins in Bacillu
J. Bacteriol. 178:4611-4619(1996)
-!- SUBCELULAR LOCATION: Cytoplasmic.
-!- INDUCTION: In response to low temperature.
-!- CAUTION: Could not be found in the genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P81095;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation updat
Cold shock protein CSI5 (11 kDa cold shock
Bacillus subtilis.
Bacteria, Firmicutes; Bacillales; Bacillace
                                                                                                                                                                    P01198;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation updat
Melanotropin alpha (Alpha-MSH).
Camelus dromedarius (Dromedary) (Arabian ca
                                                  SPECIES=C.dromedarius;
MEDLINE=75146434; PubMed=1125179;
Li C.H., Danho W.O., Chung D., Rao A.J.;
"Isolation, characterization, and amino acid melanotropins from camel pituitary glands.";
Biochemistry 14:947-952(1975).
                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camel
NCBI_TaxID=9838, 9796;
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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                                                                                                                   SEQUENCE.
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2; Conservative
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TISSUE=Pituitary;
C.H.;
and structure of
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42.9%;
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1; Mis
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Pred.
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 alpha-melanocyte-stimulating
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No.
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shock protein) (Fragment).
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n Bacillus subt
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2.8e+03;
2;
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Camelidae; Camelus
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RESULT 30
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RESULT 31
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-!- SIMILARITY: Belon.
PIR; A01464; MTCMAD.
PIR; A91785; MTHOAD.
                                                           P82239;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2001 (Rel. 42, Last annotation update)
10-OCT-2003 (Rel. 42, Last annotation update)
Salmocidin 2B (Fragment).
Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorh
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncori
NCBI TaxID=8022;
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P82238;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence up
16-OCT-2001 (Rel. 40, Last annotation
Salmocidin 2A (Fragment).
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MOD_RES 1 1 ACETYLATION
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                                                   NCBI_TaxID=8022;
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- SUBCELLULAR LOCATION: Secreted.
- TISSUE SPECIFICITY: Plasma serum.
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Pred. No.
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15-JUL-1998
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"Two-dimensional gel electrophoresis separation and N-terminal
"Two-dimensional gel electrophoresis separation and N-terminal
sequence analysis of proteins from Clostridium pasteurianum W5.";
Electrophoresis 19:802-806(1998).

Electrophoresis 19:802-806(1998).

-!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN
-!- PROTEIN IS: 5.8, ITS MW IS: 42.7 kDa.
                                           Shigella flexneri.
Bacteria; Proteoba
                                                                                                      Escherichia coli,
Escherichia coli
Escherichia coli
                                                                                                                                                                                              S4805.
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Bacteria; Firmicutes; Clo
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15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
Unknown protein CP 12 from 2D-page (Fragment).
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SEQUENCE FROM N.A.

SPECIES=E.coli; STRAIN=0157:H7 / RIMD 0509952;

SPECIES=E.coli; STRAIN=0157:H7 / RIMD 0509952;

MEDLINE=21156231; PubMed=11258796;

Hayashi T., Makino K., Ohnishi M., Kurokawa K., Ishii K., Yokoya Hayashi T., Makino K., Ohnishi M., Murata T., Tanaka M., Tobe T. Iida T., Takami H., Honda T., Sasakawa C., Ogasawara N., Yasunag Kuhara S., Shiba T., Hattori M., Shinagawa H.;

"Complete genome sequence of enterohemorrhagic Escherichia coli 0157:H7 and genomic comparison with a laboratory strain K-12.";
DNA Res. 8:11-22(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SPECIES=E.coli; STRAIN=K12 / MG1655;
MEDLINE=97426617; PubMed=9278503;
Blattner F.R., Plunkett G. III, Bloch C.A., Perna N.T., Burland V Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew G.F., Gregor J., Davis N.W., Kirkpatrick H.A., Goeden M.A., Rose D.J., Mau B., Shao Y.;
                                                                                                                                                                                                                              MEDLINE=21074935; PubMed=11206551;
Perna N.T., Plunkett G. III, Burland V., Mau B., Glasner J.D.,
Rose D.J., Mayhew G.F., Evans P.S., Gregor J., Kirkpatrick H.A.,
Posfai G., Hackett J., Klink S., Boutin A., Shao Y., Miller L.,
Grotbeck E.J., Davis N.W., Lim A., Dimalanta E.T., Potamousis K.,
Apodaca J., Anantharaman T.S., Lin J., Yen G., Schwartz D.C.,
Welch R.A., Blattner F.R.;
                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N. SPECIES=E.coli;
                                                                                                                                                                                                                                                                                                                                                                                           of uropathogenic | Proc. Natl. Acad.
                                                                                                                                                                                                                                                                                                                                                                                                     SPECIES=3.coli; STRAIN=06:H1 / CFT073 / ATCC 700928;
MEDLINE=22388234; PubMed=12471157;
Welch R.A., Burland V., Plunkett G. III, Redford P., Ro Rasko D., Buckles E.L., Liou S.-R., Boutin A., Hackett Mayhew G.F., Rose D.J., Zhou S., Schwartz D.C., Perna N Mobley H.L.T., Donnenberg M.S., Blattner F.R.;
"Extensive mosaic structure revealed by the complete general of urgathogenic Escherichia coli.";
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=80101455; PubMed=118451;
Oxender D.L., Zurawski G., Yanofsky C.;
"Attenuation in the Escherichia coli tryptophan ope secondary structure involving the tryptophan codon proc. Natl. Acad. Sci. U.S.A. 76:5524-5528(1979).
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MEDLINE=76240562; Pu
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Yanofsky C., Platt T., Crawford I.P.,
Horowitz H., van Cleemput M., Wu A.M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "The complete nucleotide sequence of the tryptophan operon Escherichia coli.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The complete genome sequence Science 277:1453-1474(1997).
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Escherichia coli.";
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., Bertrand K.,
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Tobe T.,
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EMBL; A04494; CAA00361.1; -.
EMBL; A004224; AAC74347.1; -.
EMBL; AE000224; AAN80196.1; -.
EMBL; AE0016760; AAN80196.1; -.
EMBL; AE005380; AAG56550.1; -.
EMBL; AP002556; BAB35260.1; -.
EMBL; AE015153; AAN42881.1; -.
EMBL; AE016982; AAP16766.1; -.
PIR; A03589; LFECW.
PIR; B85761; B85761.
                                                                                                                    28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
Methyl-coenzyme M reductase II gamma subunit
gamma) (Fragment).
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Tryptophan
SEQUENCE
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Cheng H.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPECIES: flexneri; STRAIN=2457T / ATCC 700930 / Serot SPECIES: flexneri; STRAIN=2457T / ATCC 700930 / Serot MEDLINE=22590274; PubMed=12704152; Wei J., Goldberg M.B., Burland V., Venkatesan M.M., De Fournier G., Mayhew G.F., Plunkett G. III, Rose D.J., Mau B., Perna N.T., Payne S.M., Runyen-Janecky L.J., 2 Schwartz D.C., Blattner F.R.; "Complete genome sequence and comparative genomics of flexneri serotype 2a strain 2457T."; Infect. Immun. 71:2775-2786(2003).
-!- FUNCTION: THIS PROTEIN IS INVOLVED IN CONTROL OF 1
                                                                                                                                                                                                                                          METTM
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MEDLINE=22272406; Pu
                                        Methanobacterium thermoautotrophicum (strain Marburg / DSM 2
Archaea; Euryarchaeota; Methanobacteria; Methanobacteriales;
Methanobacteriaceae; Methanothermobacter.
NCBI TaxID=79929;
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MEDLINE=91099370;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; A03589; LFECW.
; B85761; B85761.
; E90858; E90858.
Gene; EG11274; trpL.
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                                                                                                                                                                                                                                                                                                          FVLK---
                                                                                                                                                                                                                                                                                                                                         PKLKHYGPGW
                                                                                                                                                                                                                                                                                                                                                                                                                                       biosynthesis;
14 AA; 1723 N
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 PubMed=2269306
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                                                                                                                                                                                                                                                                                                                                                                                     25.4%;
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J., Darling
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SOURCE PARK COCCOS
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Best Local S
Matches 3
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Duve H., Johnsen A.H., Maestro J.-L., Son Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostate peptides 18:1301-1309(1997).
-:- SIMILARITY: Belongs to the allatostate Neuropeptide; Amidation.
MOD RES 8 8 AMIDATION.
          CCKN MACEU
P30369;
01-APR-1993
01-APR-1993
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ALL1 CYDPO
P82152;
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to methane and an heterodisulfide.
-!- CATALYTIC ACTIVITY: CH(3)-S-COM + H-S-HTP = CH(4)
-!- COFACTOR: Binds 2 coenzyme F430 noncovalently per
Coenzyme F430 is a yellow nickel porphinoid.
-!- PATHWAY: Methanogenesis; last step.
-!- SUBUNIT: Hexamer of two alpha, two beta, and two generates are two MCR COMPLEXES.
-!- DEVELOPMENTAL STAGE: THERE ARE TWO MCR COMPLEXES.
CONTAINS MOSTLY MCR I.
                                                                                                                                                                                                                                                                                                                                                     Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
NCBI_TaxID=82600;
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Methanogenesis; Oxidoreductase;
INIT MET 0 0
NON_TER 14 14
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                                                                                                                                                                                                               SEQÜENCE
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30-MAY-2000
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FUNCTION: Reduction of methyl-coenzyme M (2-(methylthio) ethanesulfonic acid) with 7-mercaptoheptanoylthreonine p
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                                                                                                                                                               Similarity
3; Conserv
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Similarity 75.0%;
3; Conservative
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1. 25, Last 1
1. 42, Last (CCK).

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    Last sequence up
    Last annotation

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Pred. No. :
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LATE GROWTH CELLS
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Cholecystokinin

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Best Local S
Matches 2
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MEDLINE=95009907; PubMed=7523108;

Merrick B.A., Patterson R.M., Wichter L.L., He C., Selkirk J.K.;

"Separation and sequencing of familiar and novel murine proteins
using preparative two-dimensional gel electrophoresis.";

Electrophoresis 15:735-745(1994).

-!- MISCELLANEOUS: On the 2D-gel the determined pl of this unkno
protein is: 5.2, its MW is: 50 kDa.

NON TER 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P38644;
01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of fibroblasts (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptides 9:429-431(1988).

-!- FUNCTION: This peptide hormone and the release of pancreatic ethe brain is not clear.

-!- SIMILARITY: Belongs to the gast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macropus eugenii (Tammar wallaby), and Dasyurus viverrinus (Southeastern quoll). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Metatheria; Diprotodontia; Macropodidae; Macropus. NCBI_TaxID=9315, 9279;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
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PROSITE; PS00259; GASTRIN; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Brain,
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Rodentia;
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100.0%; Pro
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                                                                                                        Score 17; Pred. No.
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Pred. No.
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1.4e+05;
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RESULT 39
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"Characterization of a new inhibitor for angiotensin converting "Characterization of a new inhibitor for angiotensin converting enzyme from the venom of Vipera aspis aspis.";

Int. J. Biochem. 22:767-771(1990).

-!- FUNCTION: This peptide both inhibits the activity of the angiotensin-converting enzyme and enhances the action of bradykinin by inhibiting the kinases that inactivate it.

It acts as an indirect hypotensive agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BPP VIPAS STANDARD; PRT; 10 AA.
P31351;
01-JUL-1993 (Rel. 26, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
Bradykinin-potentiating peptide (Angiotensin-converting enzyme inhibitor).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniati
Lepidosauria; Squamata; Scleroglossa;
Viperidae; Viperinae; Vipera.
NCBI_TaxID=8706;
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P31799;
                                                                                                   Hypotensive agent; Pyrrolidone carboxylic acid.

MOD_RES 1 1 PYRROLIDONE CARBOXYLIC ACID
SEQUENCE 10 AA; 1062 MW; 3BA827C327686773 CRC64;
                                                                                                                                                                       It acts as an in PIR; A60377; XASNPC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=92179466; PubMed=1796179;
Schoofs L., Holman G.M., Hayes T.K., Nachman R.J., de Loof A.;
"Isolation, identification and synthesis of locustamyoinhibiting
peptide (LOM-MIP), a novel biologically active neuropeptide from
Locusta migratoria.";
Regul. Pept. 36:111-119(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence upd
01-OCT-1993 (Rel. 27, Last annotation u
Locustamyoinhibiting peptide (LOM-MIP).
Locusta migratoria (Migratory locust).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vipera aspis (Aspic viper).
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Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridoidea; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=90382616; PubMed=2169439;
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     Similarity
2; Conserv
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23.9%; 5
100.0%; Pr
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100.0%; Pr/
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         Score 17; DB; Pred. No. 3.8
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387D7DD4472AB6C3
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Pred. No.
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Acridomorpha;
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RESULT 41
CAER_LITXA
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   OCCUPATION NO COCO
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Best Local Similarity
Matches 2; Conserv
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MOD_RES
SEQUENCE
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MEDLINE=20069371; PubMed=10601876;
Wabnitz P.A., Bowie J.H., Tvler " -
                                                  15-JUL-1998
15-JUL-1998
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                               Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C., Smith B.P.;
"Differences in the skin peptides of the male and female Australian tree frog Litoria splendida. The discovery of the aquatic male sex pheromone splendipherin, together with Phe8 caerulein and the antibiotic peptide caerin 1.10.";
Eur. J. Biochem. 267:269-275(2000).
-!- FUNCTION: Hypotensive neuropeptide (Probable).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          using
Rapid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Caerulein 1.2/1.274.
                                                                                                        CAER
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NCBI_TaxID=94770, 3034!
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Litoria splendida (Magnificent tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea; Hyl
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16-OCT-2001
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Litoria xanthomera (Orange-thighed frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
                                        Caerulein.
                                                                                                                                                                                                                                                                                                                                                                                    SIMILARITY:
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1306 MW;
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                                                     sequence up
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                                                                                                                                                                                                                                      Score 17;
Pred. No.
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AMIDATION.
99DBFCD37861BB5A CRC64;
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isoform 1.2 in
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3.8e+03;
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Hylidae;
   Euteleostomi;
Hylidae;
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RESULT 42
TKN1_SCYCA
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   Query Match
Best Local
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PROSITE; PSO
Tachykinin;
MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "New caerin antibacterial peptides from the skin glands of the Australian tree frog Litoria xanthomera.";
J. Pept. Sci. 3:181-185(1997).
-!- FUNCTION: Hypotensive neuropeptide.
-!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin dorsal glands.
-!- MASS SPECTROMETRY: MW=1354; METHOD=FAB.
-!- SIMILARITY: Belongs to the gastrin/cholecystokinin family.
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01-AUG-1988 (Rel.
10-OCT-2003 (Rel.
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MOD_RES 1 1
MOD_RES 4 4
MOD_RES 10 10
SEQUENCE 10 AA; 1290 MW;
                                                                                         -!- SUBCELLULAR LOCATION:
-!- SIMILARITY: Belongs to
PIR; A24867; A24867.
                                                                                                                                                                                                       Eur. J. Biochem. 214:469-474(1393).
-!- FUNCTION: Tachykinins are active peptides which excite neurons,
evoke hehavioral responses, are potent vasodilators and
                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=86192829; FubMed=2422058; Conlon J.M., Deacon C.F., C'Toole L., Thim L.; "Scyliorhinin I and II: two novel tachykinins: FEBS Lett. 200:111-116(1986).
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PROSITE; PS00259; GASTRIN; 1.
Amphibian defense peptide; Hypotensive agent; Amidation; Sulfation;
                                                                                                                                                                                                                                                             Waugh D., Wang Y., Hazon N., Balment R.J., Conlon J.M.; "Primary structures and biological activities of substance-P-related peptides from the brain of the dogfish, Scyliorhinus canicula."; Eur. J. Biochem. 214:469-474(1993).
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MEDLINE=86192829;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scyliorhinidae; Scyliorhinus.
NCBI_TaxID=7830;
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Elasmobranchii; Galeomorphii; Galeoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scyliorhinus canicula (Spotted dogfish)
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Steinborner S.T., Waugh R.J.,
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NCBI_TaxID=79697;
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                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Brain;
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                                               ; IPR002040; Tachy_Net
PS00267; TACHYKININ;
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                                                                                                                   Belongs to
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08, Last sequence 42, Last annotations
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Waugh R.J., Bowie J.H., Wallace J.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SPECTROMETRY
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                         Amidation
                                                                                                                                             Secreted.
                                                                         Neurokinin
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Pred. No.
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99DBF3837861BB5A CRC64;
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     AMIDATION
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3.8e+0
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RESULT 44
TKS2_AEDAE
TKS2_ACDAE
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01-NOV-1995
01-NOV-1995
10-OCT-2003
                    Champagne D.E., Ribeiro J.M.C.;
"Sialokinin I and II: vasodilatory tachykinins
mosquito Aedes aegypti.";
Proc. Natl. Acad. Sci. U.S.A. 91:138-142(1994).
-I- FUNCTION: Vasodilatory peptide. May activat
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Proc. Natl. Acad. Sci. U.S.A. 91:138-142(1994).
-!- FUNCTION: Vasodilatory peptide. May activat
                                                                                                                                   STRAIN=Rockefeller; TISSUE=Salivary gland;
MEDLINE=94105119; PubMed=8278354;
                                                                                                                                                                                                                          Aedes aegypti (Yellowfever mosquito).
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Aed
NCBI_TaxID=7159;
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-!- SIMILARITY: Belongs to the tacl
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Champagne D.E., Ribeiro J.M.C.;
"Sialokinin I and II: vasodilatory tachykinins
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STRAIN=Rockefeller;
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Aedes.
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RESULT 45
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OS Chlamy
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-!- MISCELLANEOUS: ON THE 2D-GEL THE E
PROTEIN IS: 6.44, ITS MW IS: 38.6
Siena-2DPAGE; P38007; -.
NON TER 10 10
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Bini L., Santucci A., Magi B.,
Comanducci M., Christianen G.,
Pallini V.;
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01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Unknown protein from 2D-page from elementary body (Fragment).
Chlamydia trachomatis.
Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.
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-!- SIMILARITY: Belongs to the tachyki
PIR; B49581; B49581.
InterPro; IPR002040; Tachy_Neurokinin.
PROSITE; PS00267; TACHYKININ; 1.
                                                                                                SEQUENCE
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MOD_RES 10 10 AMIDATION
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             Ig heavy chain CRD hypertrehalosemic adipokinetic hormo neuropeptide Led-C
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                                                                                                                                                                         hypertrehalosemic
                                                                                                                                                                                 hypertrehalosemic
                                                                                                                                                                                              hypertrehalosemic
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ALIGNMENTS

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A;Reference number: A61348; MUID:72228738; PMID:5041363
A;Accession: A61348
A;Molecule type: protein
A;Residues: 1-8 <FER1>
B;ccei-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    red pigment-concentrating hormone - northern shrimp
N;Alternate names: blanching hormone
C;Species: Pandalus borealis (northern shrimp)
C;Date: 02-Aug-1994 #sequence_revision 05-Aug-1994 #text_change 07-May-1999
C;Accession: A61348; S07139
R;Fernlund, P.; Josefsson, L.
Science 177, 173-175, 1972
A;Title: Crustacean color-change hormone: amino acid sequence and chemical and chemical sequence an
                                                                                                                                               hypertrehalosemic hormone - German cockroach
N;Alternate names: Bld-HrTH
C;Species: Blattella germanica (German cockroach)
C;Date: 03-Feb-1993 #sequence_revision 03-Feb-1993 #text_change 31-Oct-1997
C;Accession: A60421; 809137
C;Accession: A60421; 809137
R;Veenstra, J.A.; Camps, F.
Neuropeptides 15, 107-109, 1990
A;Title: Structure of the hypertrehalosemic neuropeptide of the German cockroach, A;Reference number: A60421; MUID:91179584; PMID:2080017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT
A60421
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A;Title: Structure of the red-pigment-concentrating hormone of the shrimp, Pandalus bore A;Reference number: S07139; MUID:75054965; PMID:4433569
A;Accession: S07139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; blocked amino end; hormone; neuropeptide; pyroglutami
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Modified site: amidated carboxyl end (Trp) #status experimental
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A; Residues: 'E', 2-8 < FER2>
A; Note: the amino-terminal residue form
C; Comment: This neuropeptide, isolated
A; Reference number: A60421;
A; Accession: A60421
A; Molecule type: protein
A; Residues: 1-10 < VEE>
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3; Conservative
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de, isolated from the eyestalks of the shrimp and active in pi
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Blatte

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RESULT 4

$08998

hypertrehalosemic neuropeptide Bld-HrTH - Madeira cockroach

C;Species: Leucophaea maderae (Madeira cockroach)

C;Species: Jeucophaea maderae (Madeira cockroach)

C;Species: Jeucophaea maderae (Madeira cockroach)

C;Date: 30-Jun-1992 #sequence_revision 14-Sep-1994 #text_change 24-Oct-1997

C;Accession: $08998

R;Gaede, G.; Rinehart, K.L.

Biol. Chem. Hoppe-Seyler 371, 345-354, 1990

A;Title: Primary structures of hypertrehalosaemic neuropeptides isolated freentalis and of the stick insect Extatosoma tiaratum assigned by tandem fast A;Reference number: $08995; MUID:90253659; PMID:2340112

A;Accession: $08998

A;Molecule type: protein

A;Residues: 1-10 <GAE>

C;Superfamily: adipokinetic hormone

C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental

F;10/Modified site: amidated carboxyl end (Thr) #status experimental
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A; Title: Primary structures of hypertrehalosaemic neuropeptides isolated froentalis and of the stick insect Extatosoma tiaratum assigned by tandem fast A; Reference number: S08995; MUID:90253659; PMID:2340112
A; Accession: S08997
A; Molecule type: protein
A; Residues: 1-10 <GAE>
C; Superfamily: adipokinetic hormone
C; Superfamily: adipokinetic hormone
C; Superfamily: adipokinetic hormone
C; Superfamily: adipokinetic hormone
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F; 10/Modified site: amidated carboxyl end (Thr) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Residues: 1-10 <GAE>
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic ac
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Thr) #status experimental
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Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
A;Title: Primary structures of hypertrehalosaemic neuropeptides isolated from the corpor entalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombard A;Reference number: S08995; MUID:90253659; PMID:2340112
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               Score 28; DB
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Pred. No.
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. 1.3e+02;
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C;Accession: A58620
R;Janssens, M.P.E:; Kellner, R.; Gaede, G.
Biochem. J. 302, 539-543, 1994
A;Title: A novel adipokinetic octapeptide found
A;Reference number: S55310; MUID:94379987; PMID:
                                                                                                                                                                                                 RESULT 7
A58620
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A:Moleculo 1: 555310
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A; Residues: 1-8 < JAN>
C; Superfamily: adipokinetic hormone
C; Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic ac:
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F; 8/Modified site: amidated carboxyl end (Trp) #status experimental
                     A; Reference number: A; Accession: A58620
                                                                                                                                         C;Species: Ischnura
C;Date: 28-Oct-1997
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C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Thr) #status experimental
A; Molecule type: protein
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Biochem. J. 302, 539-543, 1994
A;Title: A novel adipokinetic
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C;Species: Nauphoeta cinerea (gray cockroach)
C;Date: 31-Mar-1988 #sequence_revision 24-Oct-1997
C;Accession: A26381
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e of a hypertrehalosaemic neuropeptide
MUID:87100208; PMID:3801028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R.; Gaede,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ۱.
د.
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Pred. No. 2.8e+05;
2; Mismatches 1;
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Pred. No.
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                                         in the damselflies:8093008
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A;Residues: 1-8 <JAN>
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; hormone; neu
F;1/Modified site: pyrrolidone carboxylic acid
F;8/Modified site: amidated carboxyl end (Trp)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gonadoliberin II - American alligator
N;Alternate names: gonadotropin-releasing hormone II
C;Species: Alligator mississippiensis (American alligator)
C;Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 18-Mar-1997
C;Accession: B60066
R;Lovejoy, D.A.; Fischer, W.H.; Parker, D.B.; McRory, J.E.; Park, M.; Lance, Regul. Pept. 33, 105-116, 1991
A;Title: Primary structure of two forms of gonadotropin-releasing hormone fine A;Accession: B60066
A;Accession: B60066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: protein
A; Residues: 1-10 < LOV>
C; Superfamily: gonadoliberin
C; Keywords: amidated carboxyl end; hormone; hypothalamus; pyroglutamic acid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F; 10/Modified site: amidated carboxyl end (Gly) #status experimental
                                                                                                                                                                   A; Molecule type: protein
A; Residues: 1-10 < LOV>
A; Experimental source: brain
C; Superfamily: gonadoliberin
C; Keywords: amidated carboxyl end; brain; hormone; pyroglutamic acid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F; 10/Modified site: amidated carboxyl end (Gly) #status experimental
                                                                                                                                                                                                                                                                                                            R;Lovejoy, D.A.; Sherwood, N.M.; Fischer, W.H.; Jackson, B.C.; Rivier, J.E.; Gen. Comp. Endocrinol. 82, 152-161, 1991
A;Title: Primary structure of gonadotropin; releasing hormone from the brain CA;Reference number: A61126; MUID:91340067; PMID:1678723
A;Accession: A61126
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C;Species: Hydrolagus colliei (spotted ratfish)
C;Date: 26-May-1994 #sequence_revision 26-May-1994 #text_change 18-Mar-1997
C;Accession: A61126
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                                        KHYGPGW 11
OHWSHGW
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Pred. No. 2.8e+05;
2; Mismatches 1
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Pred. No.
                                                                                    Score 26; DB 1;
Pred. No. 2.9e+0
2; Mismatches
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2;
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gonadoliberin II - spiny dogfish N;Alternate names: gonadotropin-releasing hormone C;Species: Squalus acanthias (spiny dogfish) C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 C;Accession: B46030 R;Lovejoy, D.A.; Fischer, W.H.; Ngamvongchon, S.; Cproc. Natl. Acad. Sci. U.S.A. 89, 6373-6377, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: protein
A; Residues: 1-10 < LOV>
C; Superfamily: gonadoliberin
C; Keywords: hormone; pyroglutamic acid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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A;Title: Distinct sequence of gonadotropin-releasing hormone (GnRH) in dogfish ha;Reference number: A46030; MUID:92335300; PMID:1631133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gonadoliberin I - spiny dogfish
N;Alternate names: gonadotropin-releasing hormone
C;Species: Squalus acanthias (spiny dogfish)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 16-Dec-1998
C;Accession: A46030
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Biochem, J. 275, 671-677, 1991
A;Title: A unique charged tyrosine-containing member of the adipokinetic hor A;Title: A unique charged tyrosine-containing member of the adipokinetic hor A;Reference number: S15422; MUID:91248100; PMID:2039445
A;Accession: S15422
A;Molecule tyrotein
A;Residues: 1-8 <BIO>
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Modified site: amidated carboxyl end (Trp) #status experimental
                                                                                                                                                                                                                                                               adipokinetic hormone - cockchafer
C;Species: Melolontha melolontha (cockchafer)
C;Date: 19-Mar-1997 #sequence_revision 31-Oct-1997
C;Accession: S15422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-10 <LOV>
C;Keywords: hormone; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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3; Conserve
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42.9%;
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2.9e+02;
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us-09-720-469a-5.aug30.rpr

Best Local Similarity 50.0%; Matches 3; Conservative

Pred. No. 2.8e+05; 1; Mismatches 2;

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HYGPGW NÝSPDW 8

Matches

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7 YGPGW 11

Query Match
Best Local S
Matches 3

Similarity

35.2%;

Score 25; Pred. No.

DB 2; Le.. 2.8e+05;

Length 8; Indels

Mismatches

,;

Gaps

0

w •

Conservative

RESULT 14 A58641

adipokinetic hormone - dor beetle C; Species: Geotrupes stercorosus (dor beetle) C; Date: 28-Oct-1997 #sequence_revision 31-Oct-1997 #text_change 31-Oct-1997 C; Date: 28-Oct-1997 #sequence_revision 31-Oct-1997 #text_change 31-Oct-1997 C; Accession: A58641 #sequence_revision 31-Oct-1997 #text_change 31-Oct-1997 C; Accession: A58641 #sequence_revision 31-Oct-1997 #text_change 31-Oct-1997 #text_c the adipokinetic hormone/ red-p

A;Molecule type: protein A;Residues: 1-8 <BIO>

C;Superfamily: adipokinetic hormone C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic ac F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;8/Modified site: amidated carboxyl end (Trp) #status experimental

Query Match Best Local S Matches 3 similarity
3; Conserv Conservative 35.2%; 1; Score 25; DB 2; Pred. No. 2.8e+05; 1; Mismatches 2 'n Length 8; 0

뮍 Ś σ HYGPGW 11

RESULT 15 S21663

neuropeptide - flower beetle (Pachnoda marginata)
C;Species: Pachnoda marginata
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 24-Jul-1997
C;Accession: S21663
R;Gaede, G.; Lopata, A.; Kellner, R.; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 373, 133-142, 1992

le, G.; Lopata, A.; Kellner, R.; Rinehart, K.L. Chem. Hoppe-Seyler 373, 133-142, 1992

A; Title: Primary structures of neuropeptides isolated from ectrometry. A; Reference number: S21663; MUID:92265187; PMID:1586453 corpora cardiaca of vari

A;Accession: S21663 A;Status: preliminary A;Molecule type: protein A;Residues: 1-8 <GAE>

ద Š Query Match Best Local S Matches 3 თ Similarity 3; Conserv HYGPGW 11 NYSPDW 35.2%; ilarity 50.0%; Conservative Score 25; DB 2; Pred. No. 2.8e+05; 1; Mismatches 2

Length 8;

0;

Gaps

0

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RESULT 16 B33995

adipokinetic hormone - black horse fly
C;Species: Tabanus atratus (black horse fly)
C;Species: Tabanus atratus (black horse fly)
C;Date: 23-Mar-1990 #sequence_revision 23-Mar-1990 #text_change 31-Oct-1997
C;Accession: A33995
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Nachman, R.J.; Vogel, V.W.; Zhang, Proc. Natl. Acad. Sci. U.S.A. 86, 8161-8164, 1989
A;Title: Primary structure of two neuropeptide hormones with adipokinetic and hypotrehal A;Reference number: A33995; MUID:90046758; PMID:2813385
A;Accession: A33995
A;Accession: A33995
A;Molecule type: protein
A;Residues: 1-8 <JAF>
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamic F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status predicted
F;8/Modified site: amidated carboxyl end (Trp) #status predicted

hypotrehalosemic hormone - black horse fly C;Species: Tabanus atratus (black horse fly) C;Date: 23-Mar-1990 #sequence_revision 23-Mar-1990 #text_change 31-Oct-1997

C;Accession: B33995
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Nachman, Proc. Natl. Acad. Sci. U.S.A. 86, 8161-8164, 1989
A;Title: Primary structure of two neuropeptide hormones with A;Reference number: A33995; MUID:90046758; PMID:2813385
A;Accession: B33995 adipokinetic and hypotrehal R.J.; Vogel, V.W.;

A;Residues: 1-10 <JAF>
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamic
C;Keywords: amidated carboxylic acid (Gln) #status predicted
F;1/Modified site: amidated carboxyl end (Tyr) #status predicted A;Molecule type: protein A;Residues: 1-10 <JAF>

Matches 7 YGPGW 11 Similarity
3; Conserv Conservative 35.2%; Score 25; Pred. No. Mismatches .3e+02; 1; Length 10; 0 Gaps

0

DB 2;

밁 Ś 4 FTPGW 8

RESULT A21114

gonadoliberin - C;Species: Oncor chum salmon

C;Species: Oncorhynchus keta (chum salmon)
C;Date: 10-Aug-1990 #sequence_revision 10-Aug-1990
C;Accession: A21114 #text_change 18-Jun-1993

R;Sherwood, N.; Eiden, L.; Brownstein, M.; Spiess, J.; Rivier, J.; Vale, W. Proc. Natl. Acad. Sci. U.S.A. 80, 2794-2798, 1983
A;Title: Characterization of a teleost gonadotropin-releasing hormone. A;Reference number: A21114; MUID:83195140; PMID:6341999
A;Accession: A21114

A;Status: preliminary A;Molecule type: protein A;Residues: 1-10 <SHE>

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Matches Query Match l Similarity
3; Conserv Conservative 35.2%; Score 25; DB Pred. No. 4.3e 2; Mismatches Length 10;

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OHWSYGW KHYGPGW 11

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RESULT 18

S39030

lysyl-bradykinin - rainbow trout C;Species: Oncorhynchus mykiss (rainbow trout) C;Date: 19-May-1994 #sequence_revision 19-Apr-1996 C;Accession: S39030 R;Conlon, J.M.; Olson, K.R. FEBS Lett. 334, 75-78, 1993 #text_change 18-Aug-2000

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napin small chain S3A - Swedish turnip (fragment)
C;Species: Brassica napus var. rapifera (Swedish turnip, rutabaga)
C;Date: 19-Mar-1998 #sequence_revision 17-Apr-1998 #text_change 07-May-1999
C;Accession: S70338
R;Neumann, G.M.; Condron, R.; Thomas, I.; Polya, G.M.
Biochim. Biophys. Acta 1295, 23-33, 1996
Biochim. Biophys. Acta 1295, 23-33, 1996
A;Title: Purification and sequencing of multiple forms of Brassica napus seed napin smal A;Reference number: S70336; MUID:96283790; PMID:8679670
A;Accession: S70338
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S70338
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A;Molecule type: protein
A;Residues: 1-10 <CON>
C;Superfamily: unassigned
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                                                                                                                                                                                                                                                                                                                                                                                    Ig heavy chain DJ region (clone C344-99) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-19
C;Accession: PH1322
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B; A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1322
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Ig heavy chain CRD3 region (clone 4-106) - C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-S
C;Accession: PT0288
R;Yamada, M.; Wasserman, R.; Reichard, B.A.
J. Exp. Med. 173, 395-407, 1991
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Best Local S
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Best Local S
Matches 3
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Best Local
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;Residues: 1-14 <WAS>
;Keywords: heterotetramer; immunoglobulin
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Pred. No. 8.8e+02;
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6.3e+02;
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6.9e+02;
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                                                        #text_change 16-Aug-1996
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cytochrome-c oxidase (EC 1.9.3.1) chain VII c, hepatic - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 07-May-1999
C;Accession: S65388; S65389
C;Accession: S65388; S65389
R;Schaegger, H.; Noack, H.; Halangk, W.; Brandt, U.; von Jagow, G.
Eur. J. Biochem. 230, 235-241, 1995
Bur. J. Biochem. 230, 235-241, 1995
A;Reference number: S65372; MUID:95324529; PMID:7601105
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S77990
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A; Residues: 1-9 < YAM>
A; Experimental source: B ly
C; Keywords: heterotetramer;
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A; Residues: 1-10 <SC2>
C; Superfamily: cytochrome-c oxidase chain VIIc
C; Keywords: oxidoreductase
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A;Molecule type: protein
A;Residues: 1-10 <SCH>
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                                                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number: S77980
A; Accession: S77990
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                                                                                                                                       A,Pathway: oxidative phosphorylation; respiratory chain C;Keywords: electron transfer; membrane-associated complex; mitochondrial
                                                                                                                                                                                                                                            A;Genome:
                                                                                                                                                                                                                                                                                                                 A; Experimental source: heart; liver
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Matches 5
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er; immunoglobulin
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Sequence Database, June 1997
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Pred. No.
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Pred. No. 9.4e+02;
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Pred. No. 9.
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HYAEGPG

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R;Scarborough, R.M.; Jamieson, G.C.; Kalish, F.; Kramer, Proc. Natl. Acad. Sci. U.S.A. 81, 5575-5579, 1984
A;Title: Isolation and primary structure of two peptides A;Reference number: A49823; MUID:84298179; PMID:6591205
A;Accession: A49823
A;Molecule type: protein
A;Residues: 1-8 <SCA>
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
                                                                                                                                                                                       adipokinetic hormone I - American cockroach N;Alternate names: periplanetin CC-1 C;Species: Periplaneta americana (American cockroach) C;Date: 03-May-1994 #sequence_revision 03-May-1994 #text_change 07-May-1999 C;Accession: A49823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N;Alternate names: Pea-CAH-I
C;Species: Blatta orientalis (oriental cockroach)
C;Date: 30-Jun-1992 #sequence_revision 24-Oct-1997 #text_change 31-Oct-1997
C;Accession: S08995
R;Gaede, G.; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
A;Title: Primary structures of hypertrehalosaemic neuropeptides isolated from the entalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom A;Reference number: S08995; MUID:90253659; PMID:2340112
A;Accession: S08995
A;Molecule type: protein
A;Residues: 1-8 <GAE>
A;Note: the amino-terminal residue forms pyrrolidone carboxylic acid; therefore, C;Superfamily: adipokinetic hormone
                                                                                                                                                                                                                                                                                          A49823
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$08995

hypertrehalosemic hormone I - oriental cockroach
N;Alternate names: Pea-CAH:I
C;Species: Blatta orientalis (oriental cockroach)
C;Date: 30-Jun-1992 #sequence_revision 24-Oct-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Cator J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0302
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A; Residues: 1-11 < YAM>
A; Experimental source: B lyr
C; Keywords: heterotetramer;
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PT0302
Ig heavy chain CRD3 region (clone 5-112) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change
C;Accession: PT0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Superfamily: adipokinetic hormone; Reuropeptide; pyroglutamic acid; Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental; 8/Modified site: amidated carboxyl end (Trp) #status experimental
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ilarity 33.3%;
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corpora cardiaca; hormone; neuropeptide; pyroglutamid
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es 2;
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C;Accession: A43976
R;Gaede, G.; Rosinski, G.
Peptides 11, 455-459, 1990
A;Title: The primary structure of the hypertrehalosemic neuropeptide from te A;Reference number: A43976; MUID:90341081; PMID:2381871
A;Accession: A43976
A;Molecule type: protein
A;Residues: 1-8 <GAE>
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Modified site: amidated carboxyl end (Trp) #status experimental
hypertrehalosemic hormone - beetle (Zophobas rugipes) C;Species: Zophobas rugipes C;Date: 03-Feb-1993 #sequence_revision 03-Feb-1993 #t C;Accession: B43976 R;Gaede, G.; Rosinski, G.
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R;Gaede, G.; Kellner, R.
Peptides 10, 1287-1289, 1989
A;Title: The metabolic neuropeptides of the A;Reference number: A44960; MUID:90160053; PA;Accession: A44960
A;Molecule type: protein A;Residues: 1-8 <GAE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypertrehalosemic hormone - yellow mealworm
C;Species: Tenebrio molitor (yellow mealworm)
C;Date: 03-Feb-1993 #sequence_revision 03-Feb-1993 #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Superfamily: adipokinetic hormone
C;Keywords: blocked carboxyl end; corpora cardiaca; hormone; neuropeptide;
C;Keywords: blocked carboxyl end; corpora cardiaca; hormone; neuropeptide;
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Modified site: blocked carboxyl end (Trp) (probably amidated) #status e
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Pred. No. 2.8e+05;
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R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Heml Blochem. Biophys. Res. Commun. 124, 350-358, 1984
A;Title: Structures of two cockroach neuropeptides assigned A;Reference number: A90118; MUID:85046530; PMID:6548628
A;Accession: A05169
                                                                                                                                                                                                                                                                                                                                                                                  gonadoliberin - sea lamprey
N;Alternate names: gonadotropin releasing hormone (GnRH)
C;Species: Petromyzon marinus (sea lamprey)
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change 18-Mar-1997
C;Accession: A01412
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A;Molecule type: protein
A;Residues: 1-8 <GAE>
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide;
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Modified site: amidated carboxyl end (Trp) #status experimental
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R;Sherwood, N.M.; Sower, S.A.; Marshak, D.R.; Fraser, B.A.; Brownstein, M.J.
J. Biol. Chem. 261, 4812-4819, 1986
A;Title: Primary structure of gonadotropin-releasing hormone from lamprey brain.
A;Reference number: A01412; MUID:86168192; PMID:3514603
A;Accession: A01412
A;Molecule type: protein
A;Residues: 1-10 <SHE>
C;Comment: This hormone was isolated from the brain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neuropeptide M-I - American cockroach
C;Species: Periplaneta americana (American cockroach)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 31-Dec-1993
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C; Keywords:
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A;Title: The primary structure of the hypertrehalosemic neuropeptide
A;Reference number: A43976; MUID:90341081; PMID:2381871
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Matches
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C;Accession: A60409

R;Simmaco, M.; Severini, C.; De Biase, D.; Barra, D.; Bossa, F.; Roberts
Peptides 11, 299-304, 1990

A;Title: Six novel tachykinin- and bombesin-related peptides from the sl
A;Reference number: A60409; MUID:90287814; PMID:2356157

A;Accession: A60409

A;Molecule type: protein
A;Residues: 1-13 <SIM>
C;Superfamily: unassigned animal peptides
C;Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic of the composition of the compositi
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R;Gaede, G.; Janssens, M.P.E.
Biol. Chem. Hoppe-Seyler 375, 803-809, 1994
A;Title: Cicadas contain novel members of the AKH/RPCH:
A;Reference number: S53789; MUID:95225985; PMID:7710694
A;Accession: S53789
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A; Residues: 1-11 < VEE>
C; Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experim
F; 11/Modified site: amidated carboxyl end (Asn) #status experimental
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C; Species: Periplaneta americana (Ameri
C; Date: 07-Sep-1990 #sequence_revision
C; Accession: S05002
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C;Species: Platypleura capensis
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 07-May-1999
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R; Veenstra, J.A.

1989
FEBS Lett. 250, 231-234, 1989
FEBS Liett. 250, 231-234, 1989
FEBS Lett. 250, 231-23
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C;Keywords: blocked amino
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C;Species: Pseudophryne guentheri
C;Date: 30-Jan-1993 #sequence_revision 30-Jan-1993 #tex
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vision 09-Apr-1998 #text_change 09-Apr-1998
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Pred. No. 1.5e+03
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R;Nakano, N.; Kikutani, H.; Nishimoto, H.; Kishimoto, T.
J. Exp. Med. 173, 1091-1097, 1991
A;Title: T cell receptor V gene usage of islet beta cell-reactive A;Reference number: PT0209; MUID:91217621; PMID:1902501
A;Accession: PT0209
A;Molecule type: mRNA
A;Residues: 1-11 <NAK>
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                    RESULT 37
PT0209
                                                                                                                                                                                                                                                T-cell receptor alpha chain V-J region (4-1-L.6) - mouse C; Species: Mus musculus (house mouse)
C; Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_C; Accession: PT0209
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R;Mummert, E.; Grimm, R.; Speth, V.; Eckerskorn, C.; Schiltz, E.; Gatenby, A.A.; Schaef. Nature 363, 644-648, 1993

A;Title: A TCP1-related molecular chaperone from plants refolds phytochrome to its photo A;Reference number: S33800; MUID:93288140; PMID:8099715

A;Accession: S33802

A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chaperone, TCP1-related - oat
C;Species: Avena sativa (oat)
C;Date: 02-Dec-1993 #sequence_revision 27-Feb-1997 #text_change 17-Mar-1999
C;Accession: S33802
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S33802
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A;Note: the nucleotide sequence was submitted C;Keywords: heterotetramer; immunoglobulin
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R;Huber, C.; Klobeck, H.G.; Zachau, H.G.
Eur. J. Immunol. 22, 1561-1565, 1992
A;Title: Ongoing V(kappa)-J(kappa) recombination after formation of a productive A;Reference number: S23637; MUID:92289816; PMID:1601042
A;Accession: S23640
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A; Residues:
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A; Residues: 1-13 < HUB>
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C;Species: Homo sapiens (man)
C;Date: 19-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 16-Aug-1996
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Pred. No. 1.8e+03;
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Pred. No. 1.9e+03;
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                                                                                                                       C;Species: Gryllus bimaculatus (two-spotted C;Date: 30-Jun-1989 #sequence_revision 24-C C;Accession: A28004 R;Gaede, G.; Rinehart, K.L. Biochem. Biophys. Res. Commun. 149, 908-914 A;Title: Primary sequence analysis by fast A;Reference number: A28004; MUID:88106553; A;Accession: A28004
A;Note: the amino-terminal residue forms pyrrolidone carboxylic acid; therefore, we C;Superfamily: adipokinetic hormone C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglut F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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                                                                                    A; Molecule type: protein A; Residues: 1-8 < GAE>
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AZ8004

Adipokinetic hormone G - two-spotted cricket

N;Alternate names: AKH-G

C;Species: Gryllus bimaculatus (two-spotted cricket)

C:Date: 30-Jun-1989 #sequence_revision 24-Oct-1997 #text_change
                                                                                                                                                                                                                                                                                                                                            R;Schwager, J.; Grossberger, D.; du Pasquier, L. EMBO J. 7, 2409-2415, 1988
A;Title: Organization and rearrangement of immunoglobulin M genes A;Reference number: S01158; MUID:89052653; PMID:2903824
A;Accession: S03530
                                                                                                                                                                                                                                                                                                                                                                                                                                A;Accession: PH1593
A;Molecule type: DNA
A;Residues: 1-13 <LEV>
A;Experimental source: bone
C;Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                            A; Cross-references: EMBL: X14918;
                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-14 <SCH>
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A;Title: Molecular characterization of transgene-induced A;Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig H chain V-D-J region (wild-type clone 144) - mon C;Species: Mus musculus (house mouse) C;Date: 02-Jun-1994 #sequence_revision 02-Jun-1994
                                                                                                                                                                                                                       Matches
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Best Local :
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908-914, 1987
by fast atom bombardment mass spectrometry of a 106553; PMID:3426616

peptio

neuropeptide; pyroglutamic experimental

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RESULT 43
PT0368
Ig gamma c
C; Species:
C; Date: 31
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S11545
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A;Reference number: S11545; MUID:90351345; PMID:2386478

A;Recession: S11545

A;Molecule type: protein

A;Residues: 1-8 <GAE>
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamic
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;8/Modified site: amidated carboxyl end (Trp) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                     adipokinetic hormone - nestling-sucking blowfly C;Species: Protophormia terraenovae (nestling-sucking blowfly) C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change C;Accession: S11545 R;Gaede, G.; Wilps, H.; Kellner, R. Biochem. J. 269, 309-313, 1990 A;Title: Isolation and structure of a novel charged member of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: protein
A; Residues: 1-8 <BIO>
C; Comment: This peptide has both adipokinetic and hypertrehalosemic activities.
C; Superfamily: adipokinetic hormone
C; Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamid
F; 1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F; 8/Modified site: amidated carboxyl end (Trp) #status experimental
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Biol. Chem. Hoppe-Seyler 371, 475-483, 1990
A;Title: The putative ancestral peptide of the adipokinetic/red-pigment-concentrating A;Reference number: S10596; MUID:90359055; PMID:2390213
A;Accession: S10596
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 g gamma chain C region (gamma-1) - hum
;Species: Homo sapiens (man)
;Date: 31-Mar-1992 #sequence_revision
;Accession: PT0368
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Pred. No. 2.8e+05;
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                   #text_change 16-Aug-1996
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R;Mu, M.; Nie, Z.Q.; Yang, J.
R;Mu, M.; Nie, Z.S.; Yang, J.
Plant Cell 1, 551-557, 1989
Plant Cell 1, 551-557, 1989
A;Title: The 18-KD protein that binds to the chloroplast
A;Title: The number: PQ0152; MUID:92404720; PMID:2562513
                                                                                                                                                                                                                                                                                                                          18K iron-sulfur protein - Chlamydomonas reinhardtii chloroplast N;Alternate names: frxB homolog C;Species: chloroplast Chlamydomonas reinhardtii C;Species: chloroplast Chlamydomonas reinhardtii C;Date: 23-Nov-1991 #sequence_revision 23-Nov-1991 #text_change
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A; Residues: 1-8 <MIL>
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2; Mismatches 1
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NADH2 dehydrogenase (EC 1.6.99.3) 26K chain - fava bean mitochondrion (fragment)
N;Alternate names: complex I 26K chain; NADH-ubiquinone reductase 26K chain
C;Species: mitochondrion Vicia faba (fava bean)
C;Date: 03-May-1994 #sequence_revision 07-Oct-1994 #text_change 03-Jun-2002
C;Accession: PQ0786
R;Leterme, S.; Boutry, M.
Plant Physiol. 102, 435-443, 1993
A;Title: Purification and preliminary characterization of mitochondrial complex I (NADH: A;Accession: PQ0786
A;Accession: PQ0786
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Mol. Immunol. 28, 753-761, 1991
A;Title: Early occurrence of immunoglobulin isotype switching in human fetal liver.
A;Reference number: PT0368; MUID:91312348; PMID:1906981
A;Accession: PT0368
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;;Comment: This enzyme catalyzes electron transfer from endogenous
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                                                                                                                                                                                                                                                                                   replicative
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Search completed: August 30, 2004, 10:58:50 Job time : 4.30743 secs

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Minimum DB
Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Post-processing:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   on:
            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                seq
geq
                                                                                                                                                                                                                                                                                                                                                                                                              Published Applications AA:*

1: /cgn2 6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

2: /cgn2 6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

3: /cgn2 6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

4: /cgn2 6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

6: /cgn2 6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

7: /cgn2 6/ptodata/1/pubpaa/US08 NEW PUB.pep:*

8: /cgn2 6/ptodata/1/pubpaa/US08 PUBCOMB.pep:*

9: /cgn2 6/ptodata/1/pubpaa/US08 PUBCOMB.pep:*

10: /cgn2 6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

11: /cgn2 6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

12: /cgn2 6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

13: /cgn2 6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

14: /cgn2 6/ptodata/1/pubpaa/US10A PUBCOMB.pep:*

15: /cgn2 6/ptodata/1/pubpaa/US10B PUBCOMB.pep:*

16: /cgn2 6/ptodata/1/pubpaa/US10B PUBCOMB.pep:*

16: /cgn2 6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

17: /cgn2 6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

18: /cgn2 6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

19: /cgn2 6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

16: /cgn2 6/ptodata/1/pubpaa/US10 NEW PUB.pep:*

17: /cgn2 6/ptodata/1/pubpaa/US10 NEW PUB.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1297172 segs, 314612898 residues
            Copyright
                                                                                                                                                                                                                                                                        Length
            2004, 10:49:45; Search time 13.4527 Seconds (without alignments) 257.253 Million cell updates/sec
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US-10-394-980-52
US-10-386-612-168
US-09-849-092-2
US-09-849-092-2
US-09-880-748-2870
US-09-880-748-2870
US-10-169-10-23
3 US-10-169-169-6
US-10-239-321A-52
6 US-10-450-036A-95
US-09-910-552-37
US-09-910-552-37
US-10-643-465-37
US-10-072-419-2
US-10-013-312-41
US-10-013-312-41
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  Sequence 52, Appl Sequence 168, Appl Sequence 2, Appli Sequence 99, Appl Sequence 2870, Appl Sequence 23, Appl Sequence 23, Appl Sequence 52, Appl Sequence 52, Appl Sequence 37, Appl Sequence 37, Appl Sequence 37, Appl Sequence 2, Appl Sequence 235, Appl Sequence 235, Appl
                                                                                                                                                                                                                                                                      Description
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| 26 | 26 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | 27 | |
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| -10-062-710-16 | US-10-182-252A-200 | -10-286-457-13 | -10-723-434-13 | -10-723-434-11 | -10-360-101-15 | -10-360-101-15 | -10-360-101-15 | -10-013-312-240 | -10-013-312-233 | -10-013-312-232 | -10-013-312-225 | -10-013-312-22 | -10-013-312-211 | -10-013-312-209 | -10-013-312-68 | -10-013-312-66 | -10-013-312-49 | -10-013-312-38 | -10-013-312-25 | -10-013-312-18 | -10-013-312-5 | -10-013-312-197 | -10-013-312-167 | -10-013-312-161 | -10-013-312-155 | -10-013-312-1 | -10-013-312-140 | -10-013-312-60 | -10-013-312-51 | |
| equence 16 | Sequence 200, App | equence 132, | equence 132, | equence 113, | equence 157, | equence 156, | equence 155, | equence 2407, | equence 2339, | equence 2321, | equence 2250, | equence 2217, | equence 2114, | equence 2094, | equence 686, | equence 664, | equence 497, | equence 384, | equence 254, | equence 183, | equence 52, I | equence 1976, | equence 1672, | equence 1612, | equence 1554, | equence 1464, | equence 1404, | equence 602, | equence 512, | |

ALIGNMENTS

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FILLE REFERENCE: VBV-001
FILE REFERENCE: VBV-001
CURRENT APPLICATION NUMBER: US/10/394,980
CURRENT FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: PCT/EP02/03368
PRIOR FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: US60/278,171
PRIOR FILING DATE: 2001-03-22
PRIOR APPLICATION NUMBER: US60/318,749
PRIOR FILING DATE: 2001-09-12
PRIOR APPLICATION NUMBER: US60/318,749
PRIOR APPLICATION NUMBER: US60/323,999
PRIOR FILING DATE: 2001-09-20
NUMBER OF SEQ ID NOS: 473
SOFTWARE: PATENTIN VETSION 3.1
SEQ ID NO 52
LENGTH: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1

US-10-394-980-52

; Sequence 52, Application US/10394980
; Publication No. US20040005633A1
; GENERAL INFORMATION:
; APPLICANT: Vandekerckhove, Joel
; APPLICANT: Gevaert, Kris
                                                                                                                         ; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: MISC_FEATURE
; OTHER INFORMATION: part of CYPB_HUMAN (peptidyl-prolyl cis-trans isomerase
US-10-394-980-52
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                                              Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Gevaert, Kris
TITLE OF INVENTION: QUANTITATIVE PROTEOME ANALYSIS, AP
                                                                                                                                                                                                                                 LENGTH: 14
TYPE: PRT
   6 HYGPGW 11
                                              Similarity 6; Conserv
                                              Conservative
                                                                                         63.4%;
                                              Score 45; DB 15;
;; Pred. No. 1.5;
0; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 172
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/09849092 Publication No. US20030032169A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: MYRISTATE LOCATION: (1)...(0) NAME/KEY: AMIDATION LOCATION: (0)...(13) OTHER INFORMATION: TrkB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Ben-Sasson, Shmuel A.
TITLE OF INVENTION: Short Peptides which Selectively
TITLE OF INVENTION: Modulate the Activity of Protein Kinases
FILE REFERENCE: 1242.1029-000 (CMCC-679)
CURRENT APPLICATION NUMBER: US/10/038,612
CURRENT FILING DATE: 2002-01-08
PRIOR APPLICATION NUMBER: US 09/161,094
PRIOR APPLICATION NUMBER: US 09/161,094
PRIOR FILING DATE: 1202-09-25
PRIOR FILING DATE: 108-09-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Equence 168, A blication No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
PRIOR APPLICATION NUMBER: 08/375,962
FILING DATE: 20-January-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/258,418
                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/729,594
                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                       SOFTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/849,092
                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 144 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
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                                                                                                                                                                                                                                                                                       COMPUTER: IBM
OPERATING SYSTEM:
                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                           STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 NEKTKHYGP 9
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1: 805 Third Avenue
New York City
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                                                                                                                                                                                                                                                                                                                                                                                                             New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eriksson, Ulf; Simon, Andras; Romert, Anna
/ENTION: ISOLATED NUCLEIC ACID MOLECULE WHICH
/ENTION: CODES FOR A 32 KDA PROTEIN HAVING 11-CIS RETINOL DEHYDROGENASE
/ENTION: ACTIVITY, AND WHICH ASSOCIATES WITH P63, A PORTION OF A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US20020160478A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2,
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                                                                                                                                                                                                                                                           US-09-880-748-2870
       FILE REFERENCE: PF523

CURRENT APPLICATION NUMBER: US/09/880,748

CURRENT FILING DATE: 2001-06-15

PRIOR APPLICATION NUMBER: 60/212,210

PRIOR FILING DATE: 2000-06-15

PRIOR APPLICATION NUMBER: 60/240,816

PRIOR APPLICATION NUMBER: 60/240,816

PRIOR APPLICATION NUMBER: 60/276,248

PRIOR APPLICATION NUMBER: 60/276,248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 80.0
                                                                                                                                                                                                   Publication No. US20
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 99
LENGTH: 11
                                                                                                                                                                                                                      Sequence 2870, Application US/09880748 Publication No. US20030059937A1
                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/791,551
CURRENT FILING DATE: 2001-02-26
PRIOR APPLICATION NUMBER: 60/185,390
PRIOR FILING DATE: 2000-02-28
PRIOR APPLICATION NUMBER: 60/233,625
PRIOR APPLICATION NUMBER: 60/233,625
PRIOR FILING DATE: 2000-09-18
NUMBER OF SEQ ID NOS: 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENERAL INFORMATION:
APPLICANT: KLOETZER, WILLIAM S.
APPLICANT: HANNA, NABIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: METHOD FOR PREPARING ANTI-MIF ANTIBODIES FILE REFERENCE: 037003/0277869
                                                                                                                                                               APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: LUTELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                      2 HYGSSW 7
                                                                                                                                                                                                                                                                                                                                                                           6 HYGPGW 11
                                                                                                                                                                                                                                                                                                                                                                                                             4; Conservative
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No. US20030235584A1
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1-03-
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                                                                                                                                                                                                                                                                                                                                                                                                             Score 31; DB 11; Length 11; Pred. No. 2.4e+02; O; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         b.
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; PRIOR FILING DATE: 2001-03-21
; PRIOR APPLICATION NUMBER: 60/29
; PRIOR FILING DATE: 2001-05-25
; NUMBER OF SEQ ID NOS: 3239
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2870
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US-10-293-418-2870
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US-09-880-748-2870
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US-10-169-710-23
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PRIOR APPLICATION NUMBER: 60/340,817
PRIOR APPLICATION NUMBER: 09/880,748
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/276,248
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Best Local Similarity 100.
                                                                                                                Sequence 23, Application US/10169710 Publication No. US20030181362A1 GENERAL INFORMATION:
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Best Local
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CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT FILING DATE: 2002-11-27
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PRIOR FILING DATE: 2001-11-16
                                                       APPLICANT: INNOGENETICS
TITLE OF INVENTION: HUMAN AND PARASITE ORPHAN RECEPTOR PROTEINS
FILE REFERENCE: 2551-79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/212,210
CURRENT APPLICATION NUMBER: US/10/169,710 CURRENT FILING DATE: 2002-07-09 PRIOR APPLICATION NUMBER: PCT/GB01/00085
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TLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/240,816
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4; Conservative
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o. US20030223996A1
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100.0%; Pred. No. 1.2e+0
tive 0; Mismatches
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US-10-239-321A-52

Sequence 52, Application US/10239321A Publication No. US20040116336A1 GENERAL INFORMATION:

APPLICANT: KITADA, Chieko APPLICANT: NISHIZAWA, Naoki APPLICANT: HINUMA, Shuji

Chieko

CURRENT APPLICATION NUMBER: US/10/239,321A CURRENT FILING DATE: 2002-09-19 PRIOR APPLICATION NUMBER: PCT/JP01/02278 PRIOR FILING DATE: 2001-03-22 PRIOR APPLICATION NUMBER: JF 12-87114

2001-03-22 NTMBER: JP 12-87114

APPLICANT: HOSOYA, Masaki
TITLE OF INVENTION: Peptide derivative
FILE REFERENCE: 2702 USOP

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; TYPE: PRT ; ORGANISM: shistosoma haematobium US-10-169-710-23
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                                                                                                                            ; LOCATION: (1)...(0)
; NAME/KEY: AMIDATION
; LOCATION: (0)...(13)
; OTHER INFORMATION: TrkB
US-10-038-612-169
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TITLE OF INVENTION: Short Peptides Which Selectively
TITLE OF INVENTION: Modulate the Activity of Protein Kinases
FILE REFERENCE: 1242.1029-000 (CMCC-679)
CURRENT APPLICATION NUMBER: US/10/038,612
CURRENT FILING DATE: 2002-01-08
PRIOR APPLICATION NUMBER: US 09/161,094
PRIOR FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 172
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 169
TENDETH 12
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 23
LENGTH: 11
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Best Local Similarity
                                                                             Query Match
Best Local
                                                             Matches
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PRIOR APPLICATION NUMBER: GB0000474.7
PRIOR FILING DATE: 2000-01-10
                                                                                                                                                                                             NAME/KEY: MYRISTATE
                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                          ENGTH:
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2 FKLRAHGP 9
                            2 FKLKHYGP
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                                                            Similarity 5; Conserv
                                                       42.3%;
milarity 62.5%;
Conservative
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p. US20020160478A1
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Pred. No.
                                                                               Score 30; DB 13;
Pred. No. 4.1e+02;
                                                               Mismatches
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3.5e+02;
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US-09-910-552-37; Sequence 37, Application US/09910552; Publication No. US20020197260A1; GENERAL INFORMATION:
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 APPLICANT: Granoff, Dan M.
APPLICANT: Moe, Gregory R.
TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT DEFINE UNIQUE
TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE PREPARATION OF VACCINE
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 1238.002
CURRENT APPLICATION NUMBER: US/09/910,552
CURRENT FILING DATE: 2001-07-23
CURRENT FILING DATE: 2001-07-23
CURRENT FILING DATE: 2001-07-23
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Best Local :
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LENGTH: 13
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APPLICANT: TANHA, JASMID
APPLICANT: NARANG, SARAM
APPLICANT: STANIMIROVIC, DANICA
TITLE OF INVENTION: SINGLE-DOMAIN BRAIN TARGETING ANTIBODY FRAGMENTS
TITLE OF INVENTION: DERIVED FROM LLAMA ANTIBODIES
FILE REFERENCE: 1339-25
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PRIOR FILING DATE: 2000-09-19
NUMBER OF SEQ ID NOS: 72
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OTHER INFORMATION: Xaa means
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ORGANISM: Artificial
FEATURE:
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APPLICATION NUMBER:
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55.6%;
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57.1%;
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Pred. No. 4.
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Pred. No. 4.1e+02;
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RESULT 13
US-10-072-419-2
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US-10-643-465-37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Pa
SEQ ID NO 37
LENGTH: 8
                                                     CURRENT APPLICATION NUMBER: US/10/072,419
CURRENT FILING DATE: 2002-02-07
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2
                                                                                                                                                                                                                                  Sequence 2, Application US/10072419 Publication No. US20030162717A1 GENERAL INFORMATION:
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Best Local Similarity
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Best Local
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NUMBER OF SEQ ID NOS: 68
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 37
                                                                                                                                                        APPLICANT: Schacter, Bernice
APPLICANT: Schacter, Lee
TITLE OF INVENTION: Compositions and Methods for Promoting Lipid Mobilization in Human
FILE REFERENCE: 10739-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Granoff, Dan M.
APPLICANT: Moe, Gregory R.
TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT DEFINE UNIQUE
TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE PREPARATION OF VACCINE
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 1238.002
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CURRENT FILING DATE: 2003-08-19
PRIOR APPLICATION NUMBER: US/08/925,002
PRIOR FILING DATE: 1997-08-27
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ORGANISM: Artificial Sequence
ORGANISM: Gromphadorhina portentosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: sequence from OTHER INFORMATION: a phage display peptide library
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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2 NFKVRH 7
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No. US20040077840A1
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Pred. No. 1.2e+06;
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Pred. No. 1.2e+0
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APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: OF INVENTION: 193P161B USEFUL IN TREATMENT AND DETECTION OF CANCER
TITLE OF INVENTION: 193P161B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 41
LENGTH: 9
TYPE: DET
                                                                                                                                                               Sequence 235, Application US/10013312
; Publication No. US20030223990A1
; GENERAL INFORMATION:
APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 19391E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
; NUMBER OF SEQ ID NOS: 3005
; SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 235
; LENGTH: 9
TYPE: DET
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US-10-013-312-41
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US-10-013-312-235
                                                                                                              ; TYPE: PRT; ORGANISM: Homo sapiens US-10-013-312-235
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Best Local Similarity 50.0%;
Matches 3; Conservative
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Publication No.
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Best Local Similarity
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                                                     Query Match
Best Local Similarity
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3 KLKHYG 8
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                                     Conservative
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                                                     38.0%;
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; Pred. No. 6.9e
2; Mismatches
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                                Score 27; DB 12; Length 9; Pred. No. 1.2e+06; 2; Mismatches 0; Indels
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RESULT 18 US-10-013-312-1404 ; Sequence 1404, Application US/10013312

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Sequence 602, Application US/10013312
; Sequence 602, Application US/10013312
; General INFORMATION:
; General INFORMATION:
; APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
; APPLICANT: HUBERT, RENE
; APPLICANT: HUBERT, RENE
; APPLICANT: GE, WANGMAO
; APPLICANT: JAKOBOVITZ, AYA
; APPLICANT: JAKOBOVITZ, AYA
; TITLE OF INVENTION: 103P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCI
; FILE REFERENCE: 51158-20063.00
; CURRENT EPPLICATION NUMBER: US/10/013,312
; CURRENT FILING DATE: 2002-05-30
; NUMBER OF SEQ ID NOS: 3005
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 602
; LENGTH: 9
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APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 512
LENGTH: 9
TYPE: DET
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US-10-013-312-512
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US-10-013-312-602
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-013-312-602
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                                                                   Query Match 38.0
Best Local Similarity 66.7
Matches 4; Conservative
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Best Local Similarity
3 KLKHYG 8
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3 KLEHFG 8
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                                                                        Score 27; DB 12;
Pred. No. 1.2e+06;
2; Mismatches 0
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Pred. No. 1.2e+06;
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RESULT 20
US-10-013-312-1554
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                                                                          Sequence 1554, Application US/10013312
Publication No. US20030223990A1
GENERAL INFORMATION:
APPLICANT: RAITANO, ARTHUR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:

APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GB, WANGMAO
APPLICANT: GB, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
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Best Local !
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LENGTH: 9
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Publication No. US20030223990A1
                                       APPLICANT:
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APPLICANT: JAKOBOVITZ, AYA

TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED

TITLE OF INVENTION: 193PLE1B USEFUL IN TREATMENT AND DETECTION OF CANCER

FILE REFERENCE: 51158-20063.00

CURRENT APPLICATION NUMBER: US/10/013,312

CURRENT FILING DATE: 2002-05-30

NUMBER OF SEQ ID NOS: 3005

SOFTWARE: Patentin Ver. 2.1
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APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, I
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 9
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                CHALLITA-EID, PIA
FARIS, MARY
HUBERT, RENE
GE, WANGMAO
                                                                                                                                                                                                                                                                                                                   38.0%; Score 27; DB 66.7%; Pred. No. 1.20 tive 2; Mismatches
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66.7%;
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Pred. No. 1.2e+06;
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GENERAL INFORMATION:
APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCE
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1612
LENGTH: 9
TYPE: PRT
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 GENERAL INFORMATION:

APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-BID, PIA
APPLICANT: CHALLITA-BID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1554
                                                                                                                                                                                                                                                                                      Sequence 1672, Application US/10013312
Publication No. US20030223990A1
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Best Local (
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Best Local
NUMBER OF SEQ ID NOS: 3005
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Local Similarity 66.7%;
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66.7%; Pred. No. 1.
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Pred. No. 1.2e+06;
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Length 10,

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Sequence 52, Application US/10013312

; Publication No. US20030223990A1

; GENERAL INFORMATION:
   APPLICANT: RAITANO, ARTHUR
   APPLICANT: CHALLITA-EID, PIA
   APPLICANT: CHALLITA-EID, PIA
   APPLICANT: HUBERT, RENE
   APPLICANT: JAKOBOVITZ, AYA
   TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
   TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
   FILE REFERENCE: 51158-20063.00
   CURRENT APPLICATION NUMBER: US/10/013,312
   CURRENT FILING DATE: 2002-05-30
   NUMBER OF SEQ ID NOS: 3005
   SOFTWARE: PatentIn Ver. 2.1
   SEQ ID NO 52
   LENGTH: 10
   TYPE: PRT
   ORGANISM: Homo sapiens

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US-10-013-312-1976
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APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 19391E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1976
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Best Local Similarity 66.7%;
Matches 4; Conservative
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SEQ ID NO 1672
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Best Local Similarity
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TYPE: PRT
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Pred. No. 1.
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Pred. No. 1.
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1.2e+06;
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APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193PIEIB USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 254
LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
US-10-013-312-254
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-013-312-183
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US-10-013-312-254
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APPLICANT: CHALLTA-SID, PIA
APPLICANT: FARIS, MARY
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 183
                                      Query Match
Best Local Similarity
Matches 4; Conser
                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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Best Local (
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Best Local Similarity
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Similarity 66.7%;
4; Conservative
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    кткнус в
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                                         Conservative
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Pred. No. 1e+03;
2; Mismatches 0
                                      Score 27; DB 12; Length 10; Pred. No. 1e+03; 2; Mismatches 0; Indels
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Pred. No. 1e+03;
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APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA.
APPLICANT: CHALLITA-EID, PIA.
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTI
ITILE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 384
; Sequence 664, Application US/10013312; Publication No. US20030223990A1; GENERAL INFORMATION:
                                                                    RESULT 29
US-10-013-312-664
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 497
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o._US20030223990A1
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b. US20030223990A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 686
                                                                                                                               Sequence 2094, Application US/10013312 Publication No. US20030223990A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 664
LENGTH: 10
TYPE: PRT
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Best Local Similarity
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APPLICANT: CHALLITA-EID, I
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SECULD NOT 201
                                                    APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, I
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
                                                                                                                                                                                                                                                                                                                                      Local Similarity
les 4; Conserv
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Pred. No. 1e+03;
2; Mismatches 0; Indels
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Sequence 2114, Application US/10013312

Publication No. US20030223990A1

GENERAL INFORMATION:
APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2114
LENGTH: 10
Sequence 2217, Application US/10013312

Publication No. US20030223990A1

GENERAL INFORMATION:
APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
FITTLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2217
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CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2094
LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
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US-10-013-312-2217
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Best Local S
Matches 4
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Best Local Similarity
Matches 4; Conserv
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4; Conserva
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66.7%;
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Pred. No.
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Pred. No.
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GENERAL INFORMATION:
APPLICANT: RAITANO, ARTHUR
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCI
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2250
LENGTH: 10
TYPE: PRT
GRANISM: Homo sapiens
US-10-013-312-2250
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-013-312-2217
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US-10-013-312-2321
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                                                  ; TYPE: PRT; ORGANISM: Homo sapiens US-10-013-312-2321
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US-10-013-312-2250
                                                                                                                         APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: HUBERT, RENE
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2321
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Query Match
Best Local Similarity
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Score 27;
Pred. No.
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Pred. No. 1e+03;
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Pred. No. 1e+03;
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 DB 12;
1e+03;
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                                                                         Query Match
Best Local S
Matches 4
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SEQ ID NO 2407
LENGTH: 10
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Best Local (
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                                                                                                                                                                                                                                                        APPLICANT: CHALLITY, PIA
APPLICANT: FARIS, MARY
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193PIE1B USEFUL IN TREATMENT AND DETECTION OF OF
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE
APPLICANT: GE, WANGMAO
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN ENTITLED
TITLE OF INVENTION: 193P1E1B USEFUL IN TREATMENT AND DETECTION OF CANCER
FILE REFERENCE: 51158-20063.00
CURRENT APPLICATION NUMBER: US/10/013,312
CURRENT FILING DATE: 2002-05-30
NUMBER OF SEQ ID NOS: 3005
SOFTMARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: RAITANO, ARTHUR APPLICANT: CHALLITA-EID, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
                                      3 KLKHYG 8
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KLEHFG 10
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2; Mismatches
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                                                                       Score 27; DB 12; Length 10; Pred. No. 1e+03; 2; Mismatches 0; Indels
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; FEATURE:
; OTHER INFORMATION: (Q1,A4,S5,C9)-sequence of LHRH2
US-10-360-101-156
                                                                                                                           RESULT 40
US-10-360-101-157
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US-10-360-101-156
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Best Local Similarity
"hes 3; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: (Q1,A4,S5,C8)-sequence of LHRH2 US-10-360-101-155
                                 ; Sequence 157, Applica
; Publication No. US200
; GENERAL INFORMATION:
; APPLICANT: Moll, Ger
; APPLICANT: Leenhout
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/360,101
CURRENT FILING DATE: 2003-02-07
PRIOR APPLICATION NUMBER: EP 02077060.8
PRIOR FILING DATE: 2002-05-24
NUMBER OF SEQ ID NOS: 309
SOFTWARE: PatentIn version 3.1
SEQ ID NO 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Moll, Gert N.
APPLICANT: Leenhouts, Cornelis J.
TITLE OF INVENTION: Export and modification
FILE REFERENCE: 2183-5673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 309
SOFTWARE: PatentIn version 3.1
SEQ ID NO 155
LENGTH: 10
                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                              Query Match
APPLICANT: Leenhouts, Cornelis J. TITLE OF INVENTION: Export and modification of (poly) peptide in the lantibiotic way FILE REFERENCE: 2183-5673
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CURRENT FILING DATE: 2003-02-07
PRIOR APPLICATION NUMBER: EP 02077060.8
PRIOR FILING DATE: 2002-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Leenhouts, Cornelis J. IIIILE OF INVENTION: Export and modification of (poly)peptide in the lantibiotic way FILE REFERENCE: 2183-5673
                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH: 10
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                                                                                        Application US/10360101
c. US20040009550A1
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5. US20040009550A1
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                                                                                                                                                                                                                                                                           Conservative
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                                                       Gert N.
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                                                                                                                                                                                                                                                                                             38.0%;
42.9%;
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Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                             Score 27;
Pred. No.
                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                               1e+03;
                                                                                                                                                                                                                                                                                                               DB 15; Length 10;
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RESULT 42
US-10-723-434-132
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; OTHER INFORMATION:
US-10-360-101-157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-723-434-113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: EP 02077060.8 PRIOR FILING DATE: 2002-05-24 NUMBER OF SEQ ID NOS: 309 SOFTWARE: PatentIn version 3.1 SEQ ID NO 157
                                                                      Sequence 132, Applic Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity Matches 3; Conserv
                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Publication No. US200
GENERAL INFORMATION:
APPLICANT: Zhong, Pingyu
APPLICANT: Luo, Peizhi
APPLICANT: Wang, Kevin C.
APPLICANT: Hsieh, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/284,407
PRIOR FILING DATE: 2001-04-17
PRIOR APPLICATION NUMBER: US 10/125,687
PRIOR FILING DATE: 2002-04-17
PRIOR APPLICATION NUMBER: US 10/153,176
PRIOR APPLICATION NUMBER: US 10/153,176
PRIOR FILING DATE: 2002-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Li, Yan
TITLE OF INVENTION: HUMANIZED ANTIBODIES AGAINST VASCULAR ENDOTHELIAL GROWTH FACTOR
FILE REFERENCE: 26050-709.501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Zhong, Pingyu
APPLICANT: Luo, Peizhi
APPLICANT: Wang, Kevin C.
APPLICANT: Hsieh, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 10/443,134 PRIOR FILING DATE: 2003-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/723,434 CURRENT FILING DATE: 2003-11-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/360,101
CURRENT FILING DATE: 2003-02-07
                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          ID NO 113
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                                                                                                           Application US/10723434
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D. US20040133357A1
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42.9%;
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Pred. No. 1e+03;
l; Mismatches
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Pred. No. 1e+03;
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Query Match
Best Local Similarity
                                                                                                               RESULT 44
US-10-182-252A-200
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US-10-286-457-132
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CURRENT APPLICATION NUMBER: US/10/723,434

CURRENT FILING DATE: 2003-11-26

PRIOR APPLICATION NUMBER: US 60/284,407

PRIOR FILING DATE: 2001-04-17

PRIOR APPLICATION NUMBER: US 10/125,687

PRIOR APPLICATION NUMBER: US 10/153,176

PRIOR APPLICATION NUMBER: US 10/153,176

PRIOR PILING DATE: 2002-05-20

PRIOR APPLICATION NUMBER: US 10/443,134

PRIOR APPLICATION NUMBER: US 10/443,134

PRIOR FILING DATE: 2003-05-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRICE APPLICATION NUMBER: 60/334822;
PRICE FILING DATE: 2002-11-01;
NUMBER OF SEQ ID NOS: 684
SOFTWARE: Patentin version 3.1
SEQ ID NO 132
                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 4
                                                     Sequence 200, Application US/10182252A Publication No. US20040072162A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 132, Application US/10286457
Publication No. US20030166004A1
GENERAL INFORMATION:
APPLICANT: JENO GYURIS et al.
TITLE OF INVENTION: ENDOTHELIAL-CELL BINDING PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 156
SOFTWARE: PatentIn version 3.1
SEQ ID NO 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: GPCI-P01-178
CURRENT APPLICATION NUMBER: US/10/286,457
CURRENT FILING DATE: 2002-11-01
APPLICANT: FOMSGAARD, ANDERS APPLICANT: BRUNAK, SOREN APPLICANT: BUUS, SOREN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Li, Yan
FITLE OF INVENTION: HUMANIZED ANTIBODIES AGAINST VASCULAR ENDOTHELIAL GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                                                                                                                                                               1 NFKLKHYGP 9
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4; Conserv
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                                                                                                                                                                                                                                                                     Conservative
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57.1%;
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Pred. No.
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APPLICANT: Chu, Yong-Liang
APPLICANT: Chu, Yong-Liang
APPLICANT: Qiu, Jian-Tai

TITLE OF INVENTION: Polymeric Conjugates for Delivery of

TITLE OF INVENTION: MHC-Recognized Epitopes

TITLE OF INVENTION: Via Peptide Vaccines

FILE REFERENCE: 3781-001-27

CURRENT APPLICATION NUMBER: US/10/062,710

CURRENT FILING DATE: 2002-02-05

PRIOR APPLICATION NUMBER: US 60/310,498

PRIOR FILING DATE: 2001-08-08

NUMBER OF SEQ ID NOS: 232

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 030307/0205
FILE REFERENCE: 030307/0205
CURRENT FILING DATE: 2003-04-10
PRIOR APPLICATION NUMBER: PCT/DK01/00059
PRIOR FILING DATE: 2001-01-29
PRIOR APPLICATION NUMBER: EP 00610017.6
PRIOR FILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: US 60/179,333
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 1388
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 200
Search completed: August 30, 2004, 11:04:57 Job time : 14.4527 secs
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                                                                                                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HIV B Cell Epitopes
US-10-062-710-162
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US-10-062-710-162
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                                                                                                                                                                             Query Match
Best Local Similarity 80.0
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 36.6%; Score 26; DB 12; Length 8; Best Local Similarity 80.0%; Pred. No. 1.2e+06; Matches 4; Conservative 0; Mismatches 1; Indels
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITLE OF INVENTION: HIV PEPTIDE AND NUCLEIC ACIDS ENCODING THEM FOR DIAGNOSIS AND ITLE OF INVENTION: CONTROL OF HIV INFECTIONS ILE REFERENCE: 030307/0205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 HYGPG 10
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                                                                                                                                                                                                          36.6%;
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Result
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Maximum
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Maximum Match 100%
Listing first 45 sm
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Perfect score:
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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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[length: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query
Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Issued Patents AA:*

1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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Gapop 10.0 ,
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71
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     GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd
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PCT-US93-06751-120
US-07-963-324-2
PCT-US93-06751-67
US-08-375-962B-2
US-08-562-114B-2
US-08-572-993-2
US-08-271-830-56
PCT-US93-06751-65
PCT-US93-06751-103
US-08-925-002-37
US-08-925-002-37
US-08-598-873-68
PCT-US93-06751-103
US-08-598-873-68
PCT-US93-06751-104
US-08-159-339A-391
PCT-US93-06751-104
US-08-159-339A-391
PCT-US93-06751-104
US-08-279-906A-5
PCT-US93-06688-8
US-08-318-970B-23
US-08-318-970B-23
US-08-318-970B-23
US-08-318-970B-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Search time 4.34797 Seconds (without alignments)
130.609 Million cell updates/sec
                                                            Sequence 120, Appli sequence 2, Appli sequence 56, Appli sequence 66, Appli sequence 65, Appli sequence 103, Appl sequence 103, Appl sequence 112, Appl sequence 68, Appl sequence 104, App sequence 102, App sequence 5, Appli sequence 8, Appli sequence 9, Appli sequ
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| 26 36.6 9 4 US-09-59-286-4 26 36.6 9 4 US-09-59-286-4 26 36.6 9 4 US-09-515-965A-920 26 36.6 10 1 US-08-341-219-15 26 36.6 10 1 US-08-341-219-15 26 36.6 10 1 US-08-253-030-23 26 36.6 10 1 US-08-253-030-25 26 36.6 10 2 US-08-253-030-25 26 36.6 10 2 US-08-687-559-10 26 36.6 10 3 US-08-912-314A-15 26 36.6 10 3 US-08-912-314A-16 36.6 10 4 US-09-59-286-15 36.6 10 4 US-09-59-286-15 36.6 10 4 US-09-59-286-15 36.6 10 4 US-09-59-286-15 | 6 36.6 9 4 US-09-515-965A 6 36.6 9 4 US-09-350-641C 6 36.6 9 4 US-09-350-641C 6 36.6 10 1 US-08-341-219- 6 36.6 10 1 US-08-341-219- 6 36.6 10 1 US-08-341-219- 6 36.6 10 1 US-08-341-219- 6 36.6 10 1 US-08-253-030- 6 36.6 10 1 US-08-253-030- 6 36.6 10 2 US-08-425-069- 6 36.6 10 3 US-08-912-314A 6 36.6 10 3 US-08-912-314A 6 36.6 10 3 US-08-912-314A 6 36.6 10 4 US-09-599-286- 6 36.6 10 4 US-09-401-415- | 45 | 44 | 43 | 42 | 41 | 40 | 39 | 38 | 37 | 36 | 35 | 34 | ω G | 32 | 31 | 30 | 29 | 0 |
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| 4 US-09-286-4 4 US-09-350-641C 4 US-09-350-641C 1 US-08-250-789A 1 US-08-341-219-1 1 US-08-253-030-1 1 US-08-253-030-1 1 US-08-253-030-1 1 US-08-425-069-2 2 US-08-687-559-2 2 US-08-912-314A 3 US-08-912-314A 4 US-09-401-415-4 | 4 US-09-592-286-4 4 US-09-834-784-920 4 US-09-815-965A-920 5 equence 4 US-09-350-641C-920 Sequence 1 US-08-341-219-15 US-08-341-219-15 US-08-253-030-23 US-08-253-030-25 US-08-425-069-19 US-08-317-844B-19 US-08-946-525-15 US-08-946-525-15 US-08-946-525-16 US-08-946-525-15 US-08- | თ | σ, | 'n | <u>ه</u> | 9 | 'n | 9 | <u>ه</u> | <u>ა</u> | 9 | ο, | 9 | 9 | ი | σ, | σ, | <u>ა</u> | • |
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| -09-599-206-209-315-965A -09-315-965A -09-350-641C -08-341-2199-A -08-341-2199-A -08-341-2199-A -08-425-069-208-687-553-030-208-687-5585-912-314A -08-912-314A -09-599-2386-4 | -09-59-286-4 -09-834-784-920 -09-315-965A-920 -09-350-641C-920 -09-350-641C-920 -09-350-789A-94 -08-250-789A-94 -08-341-219-15 -08-341-219-16 -08-341-219-16 -08-253-030-23 -08-253-030-25 -08-425-069-19 -08-425-069-19 -08-425-069-19 -08-912-314A-16 -08-912-314A-16 -08-912-314A-16 -09-599-286-15 -09-401-415-10 -09-401-415-10 -09-599-286-15 -09-401-415-10 | 4 | 4 | w | w | w | N | N | ш | سر | مــا | 4 | μ | μ | 4 | 4 | 4 | 4 | н |
| | drence ance drence | -401-415- | -09-599-286 | -08-912-314A-1 | 08-912-314A-1 | -08-946- | -08-317-B | -08-687-559-1 | -08-425-069-1 | -08-253-030-2 | 08-253-030-2 | -08-341-219-1 | 08-341-219-1 | 08-250-789A-9 | -09-350-641C-92 | -09-515-965A-92 | -09-834-784-92 | -09-599-286- | C C C L L C L C L L |

ALIGNMENTS

Application PC/TUS9306751

GXG Epitopes

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MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE: Rando
PCT-US93~06751-120
                                                     Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                TELEFAX: (908) 594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/06751
FILING DATE: 19930719
CLASSIFICATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REFERENCE/DOCKET NUMBER: 18614
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-4678
TELEPAX: (908) 594-4720
TELEPAX: (908) 594-4720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold TITLE OF INVENTION: Immunological Conjugates of OMPC and TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization NUMBER OF SEQUENCES: 146
CORRESPONDENCE ADDRESS:
ADDRESSEE: Merck & Co., Inc.
STREET: P.O. Box 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: NJ
COUNTRY: US
ZIP: 07065
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STREET: P.O.
CITY: Rahway
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linear
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                                                                       52.1%;
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                                                                        Score 37; DB 5; Pred. No. 6.2;
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RESULT 3
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US-07-963-324-2
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Sequence 67, Application PC/TUS9306751

GENERAL INFORMATION:
APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold
TITLE OF INVENTION: Immunological Conjugates of OMPC and
                                                                                                                                                                                                      Query Match
Best Local (
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Patent No. 5274122
GENERAL INFORMATI
                                                                                                                                                                                         Matches
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (908)594-3901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OLECULE TYPE:
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                                                                                                                                                                                                                                                                LOCATION: 2..9
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: /label= nle
OTHER INFORMATION: /note= "norleucine"
                                                                                                                                                                                                                                                                                               NAME/KEY: Cross-links LOCATION: 2..9
                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US
FILING DATE: 19921015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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ISTRATION NUMBER: 35,746
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6; Conserv
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                                                                                                                             LKHIGPG 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Leanza, William J
Tolman, Richard L
TVENTION: Acidic Derivatives of Homocysteine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (908)594-4720
(908)594-4720
70 NO: 2:
                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                    49.3%; Score 35; DB 1; Length 9; 85.7%; Pred. No. 3e+05;
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/note= "amide bond through Lys epsilon amino"
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RESULT 4
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PCT-US93-06751-67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 71.4
"hes 5; Conservative
                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/08375962B Patent No. 5731195
                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
SOFTWARE: Wordperfect (ASCII standard)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/375,962B
FILING DATE: 20-January-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes NUMBER OF SEQUENCES: 146
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Merck & Co., Inc.
                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Diskette, 3.5 inch, 144 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION: (908) 594-4678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: P.O. CITY: Rahway
                                                              OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 19930
                                                                                COMPUTER:
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                                                                                                                                                                                            ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 KLKHYGP 9
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                                                                                                                                                             New York City
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                                                                                                                                            New York
                                                                                                                                                                            805 Third Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                         SIMON, ANDRAS; HELLMAN, ULF; WERNSTEDT CHRISTER; ERIKSSON, ULF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . peptide
                                                                                IBM
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O. Box 2000
                                                                                                                                                                                              Felfe & Lynch
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Portion of a Retinol Binding
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ty, and Which Associates Wi
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CLASSIFICATION:

APPLICATION DATA: LICATION NUMBER:

08/258,418

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Best Local Similarity
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                                                                                                                         FILING DATE: 22-No. 5972646ember-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/375,962
FILING DATE: 20-January-1995
ATTORNEY/AGENT INFORMATION:
NAME: KOhli, Vineet
REGISTRATION NUMBER: 37,003
REFERENCE/DOCKET NUMBER: LUD 5372.2 CI
TELECHMONE: (212) 688-9200
TELEPHONE: (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 838-3884 INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                      TELEFAX: (212) 838-3884 INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           quence 2, Aptent No. 597
                                                                                                                                                                                                                                                                                          COMPUTER: IBM
COMPUTER: IBM
COPERATING SYSTEM: PC-DOS
SOFTWARE: Wordperfect 5.1 and ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/562,114B
APPLICATION NUMBER: 22-No. 5972646ember-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 6-October-1994
ATTORNEY/AGENT INFORMATION:
NAME: Pasqualini, Patricia
                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: New York
ZIP: 10022
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 144 kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: 1i
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REFERENCE/DOCKET NUMBER: LUD 5372.1 CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                  COPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SSEE: Felfe & Lynch
T: 805 Third Avenue
New York City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4; Conservative
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                                  amino acid
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VENTION: ISOLATED NUCLEIC ACID MOLECULE WHICH CODES FOR A

VENTION: 32 KDA PROTEIN HAVING 11-CIS RETINOL HYROGENASE

VENTION: 32 KDA PROTEIN HAVING 11-CIS RETINOL HYROGENASE
                                                        8 amino acids
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linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACTIVITY, AND WHICH ASSOCIATES WITH FOS, A PORTION OF A RETINOL BINDING PROTEIN RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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Pred. No. 3e+05;
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RESULT 7 US-08-937-993-2

Sequence 2, Application US/08937993 Patent No. 6399344 GENERAL INFORMATION:

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                                                                                     Query Match
Best Local (
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Best Local
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                                                                       Matches
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APPLICATION NUMBER: 08/562,114
FILING DATE: 22-No. 6280997ember-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/375,962
FILING DATE: 20-January-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/258,418
FILING DATE: 10-June-1994
                                                                                                                                                                                                                            TELEFAX: (212) 838-3884 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION: APPLICANT: Erikse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM
OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDDERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/C
FILING DATE: 11-October-
CLASSIFICATION: 435
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 41
CORRESPONDENCE ADDRESS:
ADDRESSEE: Felfe & Lynch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Eriksson, Ulf; Simon, Andras; Romert, Anna
TITLE OF INVENTION: ISOLATED NUCLEIC ACID MOLECULE WHICH
TITLE OF INVENTION: CODES FOR A 32 KDA PROTEIN HAVING 11-CIS RETINOL DEHYDROGENASE
TITLE OF INVENTION: ACTIVITY, AND WHICH ASSOCIATES WITH P63, A PORTION OF A
TITLE OF INVENTION: RETINOL BINDING PROTEIN RECEPTOR
                                                                                                                                                           MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                               NAME: Hanson, No. 6280997man REGISTRATION NUMBER: 30,946 REFERENCE/DOCKET NUMBER: LUD
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ZIP: 10022
                                                                                                                                                                            TOPOLOGY:
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1 YSPGW
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                                  YGPGW 11
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                                                                       Conservative
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11-October-1996
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                                                                                     43.7%;
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                                                                     Score 31; DB 3; Pred. No. 3e+05; O; Mismatches
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; Sequence 56, Appli
; Patent No. 5510332
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                 APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: LUTELECOMMUNICATION INFORMATION: TELEPHONE: (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/375,962
FILING DATE: 20-January-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/258,418
FILING DATE: 10-June-1994
ATTORNEY/AGENT INFORMATION
ATTORNEY/AGENT INFORMATION
                                                     NUMBER OF SEQUENCES: 1: CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                       TITLE OF INVENTION: A PROCESS OF INHIBITING THE BINDING OF THE TITLE OF INVENTION: INTEGRIN '4 1 TO VCAM OR FIBRONECTIN AND TITLE OF INVENTION: LINEAR PEPTIDES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/562,114
FILING DATE: 22-No. 6399344ember-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/729,594
FILING DATE: 11-October-1996
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Fulbright & Jaworski L.L.P.
STREET: 805 Third Avenue
CITY: New York City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Eriksson, Ulf; Simon, Andras; Romert, Anna
TITLE OF INVENTION: ISOLATED PROTEINS HAVING CIS-R
TITLE OF INVENTION: DEHYDROGENASE ACTIVITY
NUMBER OF SEQUENCES: 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CINT.
STATE: No.:
10022
ADDRESSEE: Dre
STREET: 180 No
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Hanson, No. 63
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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KOGAN, 11...
Ren, Kaijun
: Ren, Kaijun
: Vanderslice, Peter
'Pack, Pamela J.
A PROCESS
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                                                                                                                                                                                                                                                              Application US/08271830
                  E: Dressler, Goldsmith, Shore & Milnamow, Ltd 180 No. 5510332th Stetson, Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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September 26, 1997
ON: 530
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80.0%;
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Pred. No. 3e+05;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCT-US93-06751-66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 42.3
Best Local Similarity 100
Matches 4; Conservative
                                                  CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REFERENCE/DOCKET NUMBER: 1861
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-4678
TELEPAX: 138875
TELEX: 138825
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IELEFAX: (312)616-5460
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/06751
FILING DATE: 19930719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold
TITLE OF INVENTION: Immunological Conjugates of OMPC and
TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes
NUMBER OF SEQUENCES: 146
CORRESPONDENCE ADDRESS:
ADDRESSE: Merck & Co., Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: NO. 5510332thrup, Thomas
REGISTRATION NUMBER: 33,268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (312)616-5400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                          ADDITE F. STREET: F. STREET: F. STREET: Rahway
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ZIP: 606
                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U
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                                                                                                                        18614
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HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE: Random Epitope Library Delta
PCT-US93-06751-66
                                                                                                                 MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE: Random Epitope Library Gamma
PCT-US93-06751-93
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                                                          Query Match
Best Local S
Matches 4
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION: APPLICANT: P. Kel
                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REFERENCE/DOCKET NUMBER: 1861
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-4678
TELEFAX: (908) 594-4720
                                                                                                                                                                                                                                                  TELEX: 138825
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JS93-06751-93
                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/06751
FILING DATE: 19930719
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold
TITLE OF INVENTION: Immunological Conjugates of OMPC and
TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: P.O. CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U
ZIP: 07065
                                                                                                                                                                                                          LENGTH: 14 amino acids TYPE: amino acid STRANDEDNESS: single
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TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                             COPOLOGY:
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                                                        l Similarity
4; Conserv
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                            нхере 10
HFGPG 8
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linear
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                                                           Conservative
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NO
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80.0%;
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                                                                       Score 30;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                18614
                                                           Mismatches
                                                                                       DB 5;
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                                                          Gaps
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RESULT 11

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COUNTRY: USA
ZIP: 07065

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION NUMBER: PCT/US93/06751
FILING DATE: 19930719
CLASSIFICATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REFERENCE/DOCKET NUMBER: 18614
TELECOMMUNICATION: (908) 594-4678
TELEFAX: (908) 594-4720
138825
SEQ ID NO: 65:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/353,400
FILING DATE:
CLASSIFICATION DATA:
APPLICATION NUMBER: GB 9324819.3
FILING DATE: 03-DEC-1993
PRIOR APPLICATION NUMBER: GB 9411089.7
FILING DATE: 03-UUN-1994
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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Sequence 65, Application PC/TUS9306751

GENERAL INFORMATION:
APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold

TITLE OF INVENTION: Immunological Conjugates of OMPC and

TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes
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Best Local
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: P.O. CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Merck & Co., Inc. STREET: P.O. Box 2000
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Pred. No. 98;
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PCT-US93-06751-103
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                                                             Query Match
Best Local S
Matches 4
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GENERAL INFORMATION:
APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold
TITLE OF INVENTION: Immunological Conjugates of OMPC and
TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG
NUMBER OF SEQUENCES: 146
CORRESPONDENCE ADDRESS:
ADDRESSEE: March & Co. T...
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Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                           TELEFAX: (908) 594-472
TELEX: 138825
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REFERENCE/DOCKET NUMBER: 18614
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-4678
TELEPAX: (908) 594-4720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATENT: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
                                                                                                                                           HYPOTHETICAL: NO ANTI-SENSE: NO IMMEDIATE SOURCE: Random Epitope Library Delta
                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: PC'
FILING DATE: 19930719
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LENGTH: 14 amino acids
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                                                                                                                                                                                                                         LENGTH: 14 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Merck & Co., Inc. STREET: P.O. Box 2000
                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
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                               3 KLKHYGPG 10
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                                                                              Similarity
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                                                               Conservative
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                                                              40.8%; Score 29; DB 5; 1
50.0%; Pred. No. 1.4e+02;
tive 3; Mismatches 1
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                                                                                                                                                                                                                                                                                               103:
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Pred. No. 1.4e
1; Mismatches
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1.4e+02;
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                                                                                            Length 14;
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Sequence 37, Application US/09910552

Patent No. 6642354

GENERAL INFORMATION:

APPLICANT: Granoff, Dan M.

APPLICANT: Granoff, Dan M.

TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT DEFINE UNIQUE

TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE PREPARATION OF VACCINE

TITLE OF INVENTION: COMPOSITIONS

FILE REFERENCE: 1238.002

CURRENT APPLICATION NUMBER: US/09/910,552

CURRENT FILING DATE: 2001-07-23

PRIOR APPLICATION NUMBER: 09/494,822

PRIOR FILING DATE: 2000-01-31

NUMBER OF SEQ ID NOS: 68

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 37

LENGTH: 8
RESULT 16
US-08-747-137-122
; Sequence 122, Application US/08747137
; Patent No. 5945033
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence: sequence from ; OTHER INFORMATION: a phage display peptide library US-09-910-552-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: sequence from ; OTHER INFORMATION: a phage display peptide library US-08-925-002-37
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US-09-910-552-37
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CURRENT FILING DATE: 1997-08-27
NUMBER OF SEQ ID NOS: 68
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 37, Application US/08925002
Patent No. 6048527
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                               Query Match
Best Local Similarity
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Granoff, Dan M.
APPLICANT: Moe, Gregory R.
TITLE OF INVENTION: USE OF MONOCLONAL ANTIBODIES THAT DEFINE UNIQUE
TITLE OF INVENTION: MENINGOCOCCAL B EPITOPES IN THE PREPARATION OF V
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 8
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                               LENGTH: 8
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                            4; Conservative
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Pred. No. 3e+05;
2; Mismatches
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Pred. No. 3e+05;
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/959,560
FILING DATE: 13-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/641,720
FILING DATE: 15-JAN-1991
ATTORNEY/AGENT INFORMATION:
NAME: APPLE, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 016197-0008
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
INFORMATION FOR SEQ ID NO: 122:
Patent No. 5928884
GENERAL INFORMATION:
APPLICANT: Croce, Carlo M.
APPLICANT: Huebner, Kay
TITLE OF INVENTION: FHIT PR
TITLE OF INVENTION: METHODS
                                                                                               Sequence 68, Appr.
                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
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FILING DATE: 12-NOV-1996
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/212,546
FILING DATE: 14-MAR-1994
PRIOR APPLICATION NUMBER: US 08/069,831
APPLICATION NUMBER: US 08/069,831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Modified-site
LOCATION: 1
OTHER INFORMATION: /product= "p-Glu"
FEATURE:
NAME/KEY: Modified-site
LOCATION: 10
OTHER INFORMATION: /product= "Thr-Amide"
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: YEN, Richard C.K.
TITLE OF INVENTION: NON-CROSSLINKED PROTEIN PARTICLES FOR
TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC USE
NUMBER OF SEQUENCES: 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: Two Embarcac
CITY: San Francisco
STATE: CA
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                                                                                                                          Application US/08598873
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50.0%;
   FHIT PROTEINS METHODS BASED
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ER: 016197-000840US
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     AND NUCLEIC ACIDS AND THEREON
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RESULT 18
US-08-605-430-68
; Sequence 68, Application US/08605430
---ent No. 6242212
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 02.2...

GENERAL INFORMATION:
APPLICANT: Croce, Carlo M.
APPLICANT: Huebner, Kay
TITLE OF INVENTION: FHIT PROTE
TITLE OF INVENTION: METHODS BI
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Vei
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/605,430
FILING DATE: 22-FEB-1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Friebel, Thomas E.
REGISTRATION NUMBER: 29,258
REFERENCE/DOCKET NUMBER: 8666-005
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 869-974
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Friebel, Thomas E.
REGISTRATION NUMBER: 29,258
REFERENCE/DOCKET NUMBER: 8666-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPAX: (212) 869-9741/8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/598,873
FILING DATE: 09-FEB-1996
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ADDRESSEE: Pennie & Edmonds
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STREET: 1155 Aver
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
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STREET: 11
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1155 Avenue of the Americas
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ilarity 100.0%;
Conservative (
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RESULT 20
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Best Local S
Matches 4
                                                                                                                                                                                                                                                  TELEFAX: (908) 594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acid
STRANDEDNESS: single
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Sequence 63, Applicat
GENERAL INFORMATION:
                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REFERENCE/DOCKET NUMBER: 1861
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-4678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                        ANTI-SENSE: NO
IMMEDIATE SOURCE: Random Epitope Library Delta
                                                                                                                                                                                               MOLECULE TYPE:
HYPOTHETICAL:
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Merck & C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold
TITLE OF INVENTION: Immunological Conjugates of OMPC and
TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes
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                                                                                                                        Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                                                       (908) 594-4720
                                                                                                           Conservative
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NO
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100.0%; Pr/
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                                                                                                                        Score 28;
Pred. No.
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                                                                                                          Mismatches
                                                                                                                        2e+02;
                                                                                                                                     DB 5;
                                                                                                                                     Length 14;
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RESULT 21
US-08-159-339A-391
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GENERAL INFORMATION:
APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold
TITLE OF INVENTION: Immunological Conjugates of OMPC and
TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes
NUMBER OF SEQUENCES: 146
                                                                                                                                                                                                                                                              Patent No. 6037135
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                              Sequence 391,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: pep HYPOTHETICAL: NO ANTI-SENSE: NO IMMEDIATE SOURCE:
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS LENGTH: 14 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION: (908) 594-4678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esteban
                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: 19930 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
les 4; Conserv
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                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 14 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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6037135
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O. Box 2000
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Pred. No.
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; LENGTH: 9 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-159-339A-391
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Best Local Similarity 66.,
There 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 102, Application PC/TUS9306751

GENERAL INFORMATION:

APPLICANT: P. Keller, A. J. Conley, A.R. Shaw, B.A. Arnold

TITLE OF INVENTION: Immunological Conjugates of OMPC and

TITLE OF INVENTION: HIV-Specific Selected Principal Neutralization GXG Epitopes

NUMBER OF SEQUENCES: 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                      FILING DATE: 19930719
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Meredith, Roy D.
REGISTRATION NUMBER: 30,777
REGISTRATION NUMBER: 1861
REFERENCE/DOCKET NUMBER: 1861
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 594-4678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
FILING DATE: US 08/027,746
TELEFAX: (908) 594-4720
TELEX: 138825
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 07065
ZIP: 07065
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM CON
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: P.O. CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE: 29-NO
                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ESPONDENCE ADDRESS:
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SYSTEM: DOS
FastSEQ for Windows Version 2.0
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                                                                                                         18614
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US-08-279-906A-5
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Best Local Similarity
Matches 4; Conserv
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Patent No. 561892
                                                          Matches
                                                                        Query Match
Best Local Similarity
                                                                                                                                                                              TELEFAX: (312) 474-044
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/279,906A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: No. 5618922and, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 32028
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS LENGTH: 13 amino acid:
                                                                                                                                  TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
ATTORNEY/AGENT IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Terada, Masaki
APPLICANT: Yoneda, Yukio
TITLE OF INVENTION: NMO3;
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 6300 Sear
CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                           TELEPHONE: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
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TYPE: amino acid
STRANDEDNESS: single
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                              HYGPG 10
HIGPG
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                                                            Conservative
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; Pred. No. 2.7e
1; Mismatches
                                                          <u>,</u>
                                                         Score 26; DB 1;
Pred. No. 3e+05;
0; Mismatches
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. 2.7e+02;
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RESULT 24

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CIII.
STATE: Mab.
STATE: Mab.
COUNTRY: U.S.A.
ZIP: 02110-2804
ZIP: 021110-2804

COMPUTER READABLE FORM:
COMPUTER: ISM PS/2 Diskette, 1.44 Mb
COMPUTER: ISM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: IBM P.C. DOS (Version 3.30)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/06688

APPLICATION NUMBER: PCT/US92/06688
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US-08-318-970B-23
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PCT-US92-06688-8
                                                                                                                                                                                                                                                                                                                                     Sequence 23, Application US/08318970B
Patent No. 5589573
GENERAL INFORMATION:
APPLICANT: Hideaki HAGIWARA, et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                   TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 744,281
FILING DATE: 13 August 1992
ATTORNEY/AGENT INFORMATION:
                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: 002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NERAL INFORMATION
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TITLE OF INVENTION: MULTIPLE ANTIGEN PEPTIDES FOR USE AS HIVITLE OF INVENTION: VACCINES
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage COMPUTER: Dell System 210; Intel 80 285 Microprocessor OPERATING SYSTEM: MS DOS 3.3 SOFTWARE: Word Perfect, Version 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                         STREET:
                                                                                                                                                                                                          ADDRESSEE:
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                                                                                                                                            Virginia
                                                                                                                                                                                     3: Law Office of Sherman and Shalloway 413 N. Washington Street
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                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                             Hideaki HAGIWARA, et al.
PENTION: AMINO ACID SEQUENCES OF ANTI-IDIOTYPIC
PENTION: ANTIBODIES AGAINST ANTI-CANCER HUMAN MONOCLONAL ANTIBODY
PENTION: AND DNA BASE SEQUENCES ENCODING THOSE SEQUENCES
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RESULT 26
US-08-946-525-4
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Best Local S
Matches 3
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (703) 549-220
TELEFAX: (703) 836-0106
INFORMATION FOR SEQ ID NO: 2
                                                                               TELEPHONE: (212) 697-5995
TELEFAX: (212) 286-0854 or 286-0082
TELEX: TWX 710-581-4766
NFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                         APPLICATION NUMBER: 08/785,696
FILING DATE: January 17, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Bogosian, Elizabeth A
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                          CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch 1.44 Mb storage diskette
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: MS-DOS
                                                                                                                                                                          REGISTRATION NUMBER: 39,911
REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Word Processor (ASCII) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Richard A. Steinbe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Arye Rul
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 90 Park Avenue
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          / Match 36.6%;
Local Similarity 42.9%;
les 3; Conservative
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/946,525 FILING DATE: October 7, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A. ZIP: 10016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: L CDR3-5
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 26,588
REFERENCE/DOCKET NUMBER: S-2371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/318,970B FILING DATE: October 6, 1994 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
TOPOLOGY: linear
ADDRESSEE:
                                                                ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 QHYSTAW 8
                                             amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rubinstein, Barry R. Bloom, YON: PEPTIDE COMPOSITIONS FOR ON: PREVENTION OF HIV
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Pred. No. 3e+05;
1; Mismatches
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R THE TREATMENT AND
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